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AAM24472 ABJ19682 ABP99572

AAB39216 AAU21256

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Human bone marrowHuman bone marrowTANGO 228.

AAB82313 AAB82315 AAB82317 AAY27129 AAY27130 AAY95966 ABB10224 AAU19018

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ABG96265 AAR94894

AAB82314 ABB84668

AAB82316 ABG74786 ABP69283 AAR13251 AAW14802

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AAB07652 AAB65866 AAR94893

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AAR39741

AAW76118 AAW71252

AAW44838 AAW81440

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ICAM-R (Intercell)

Human ICAM-R prote

Human ICAM-R encod

Human ICAM-R encod

Human ICAM-R prote

Human polypeptide

Lung cancer associ

Human Polypeptide

Lung cancer associ

Human A33 antigen.

AAY82435 AAY50743 AAB50095

AAB13036

AAE34390 AAB82318 AAB58139 AAW14146

AAB65863 ABP62881

ABG12152 ABB05044

AAE14781 ABR40465

Amino acid sequenc Human A33 protein Human polypeptide Human immunoglobul Novel human diagno Human secreted pro Amino acid sequenc Mouse A33 antigen. Human TANGO 330 fo Human PROSEO protein Human TANGO 330 fo Human PROSEO

Human PRO860 polyp Novel human protei Human ECSM4 protei Amino acid sequenc Novel human diagno

AAU99419 AAB31889

AAU00501 AAY41716 AAB44272

ABU61102

AAW14158

Murine Protein #6 encodeu Human FAIL protein

AAB59829 AAE23555

ABP96857

Human expressed pr Murine nephrin pro Protein #6 encoded Sequence of human

The present invention describes an isolated antibody that binds to one of the human PRO proteins designated PRO212, PRO290, PRO31, PRO1909, PRO619, PRO1717, PRO619, PRO1909, PRO2199, PRO2199, PRO319, PRO9194, PRO1817, PRO1181, PRO1182, PRO18184, PRO2187, PRO1909, PRO2199, PRO2199, PRO3199, PRO9194, PRO2199, PRO2199, PRO2199, PRO2199, PRO2199, PRO3197, PRO1909, PRO2199, PRO2199, PRO2199, PRO2199, PRO3197, PRO1909, PRO2199, PRO2199, PRO3197, PRO1909, PRO2199, PRO2199, PRO3197, PRO1909, PRO2199, PRO2199, PRO3197, PRO1909, PRO2199, PRO3199, PRO3197, PRO1909, PRO2199, PRO3199, PRO3199,

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Roy MA;

Hillan KJ,

Goddard A, Gurney AL,

Baker KP, Wood WI;

Ashkenazi AJ,

Watanabe CK,

WPI; 2000-572270/53. N-PSDB; AAC58373.

(GETH) GENENTECH INC.

Thirty PRO polynucleotides encoding PRO polypeptides, useful in the treatment, diagnosis and prevention of cancer -

Claim 61; Fig 14; 286pp; English.

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Human; tumour; diagnosis; neoplastic disease; neoplastic cell growth; proliferation; tumourigenesis; identification; cancer; cytostatic; nootropic; neuroprotective; antiinflammatory; immunosuppressive; immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy; neuronal disorder; glial disorder; astrocytal disorder; angiogenic; hypothalamic disorder; glandular disorder; macrophagal disorder; epithelial disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; immunologic disorder.
                                                   Human PRO809 protein sequence SEQ ID NO:23.
AAB24063 standard; Protein; 265 AA.
                                                                                                                                                                                                                             99WO-US12252.
99US-0141037.
99US-0145698.
99US-0145698.
99WO-US28313.
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20-DEC-1999;
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Best Local Similarity 100.0%;
Matches 265; Conservative 0;
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98US-0088810.
98US-0088811.
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 265 AA;
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02 - 70N - 1998

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10-JUN-1998;
10-JUN-1998;
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Sequence
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The invention provides membrane-bound PRO polypeptides and polymucleotides encoding them. The PRO sequences of the invention were identified based on extracellular domain homology screening. The PRO sequences have homology with proteins including LDL receptors, TIE ligands and various enzymes. The membrane-bound proteins and receptor molecules are useful as pharmaceutical and diagnostic agents. Receptor immunoadhesins, for instance, can be used as therapeutic agents to block receptor-ligand interactions. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. The PRO encoding sequences are useful as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. PRO nucleic acid sequences will also be useful for the preparation of PRO polypeptides, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Smith V, Watanabe CK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 1409; DB 21; Length 265; 100.0%; Pred. No. 2.9e-131; cive 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Membrane-bound proteins and related nucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gurney AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            claim 12; Fig 151; 822pp; English.
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Wood WI, Yuan J;
                                                                                    98US-0096757
98US-0096766
98US-0096778
98US-00968773
98US-0096897
98US-0096895
98US-0096895
98US-0096895
98US-0096969
98US-0096969
98US-0096969
98US-0097022
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98US-0097052
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98US-0095929.
98US-0096012.
98US-0096146.
98US-0096146.
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98US-0097979-
98US-0097014-
98US-0098515-
98US-0100634-
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Best Local Similarity 100.
Matches 265; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GETH ) GENENTECH INC.
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18-AUG-1998;
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18-AUG-1998;
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17-AUG-1998;
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17-AUG-1998
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NESFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKWEDWQGPLESPILALPLYRSTR 240
                                                                                             NFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKWEDWQGPLESPILALPLYRSTR 240
                                                                      121 LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA 180
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      SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWE
                         SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWE
                                                  LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA
                                                                                                                                                                                                                                                                                                           Human; secreted and transmembrane protein; PRO; cytostatic;
cell death; cancer; chromosomal mapping; gene mapping; tissue typing;
diagnostic assay.
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Wood WI;
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                                                                                                                                                                                                                                                                                          Human PRO809 (UNQ464) protein sequence SEQ ID NO:223.
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                                                                                                                                           241 RLSEEFFGGFRIGNGEVRGRKAAAM 265
                                                                                                                                                         RISEEEFGGFRIGNGEVRGRKAAAM 265
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                                                                                                                                                                                                                       AAB65214 standard; Protein; 265 AA.
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99US-0143048.
99US-0144758.
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99US-0146222.
99US-0149396.
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99US-0158663
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Grimaldi CJ, Gurney
Roy MA, Stewart TA,
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15-SEP-1999;
08-OCT-1999;
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11-FEB-2000;
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15-MAR-2000;
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Smith V, Stephan JF,
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2000US-220664P.
2000US-220666P.
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2000WO-US20710.
2000WO-US23522.
2000WO-US23328.
2000US-00000P.
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2000WS-253646P.
2000WO-US32678.
2000WO-US34256.
2001WO-US34956.
2001WO-US06520.
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2000US-220605P.
2000US-220607P.
2000US-220624P.
               2001WO-US21066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    as lung cancer,
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                                                                          25-JJL-2000; 25-JJL-2000; 25-JJL-2000; 25-JJL-2000; 25-JJL-2000; 26-JJL-2000; 28-JJL-2000; 28-JJL-2000; 2
                                                                                                                                                                      23-AUG-2000; 24-AUG-2000; 215-SEB-2000; 210-NOV-2000; 28-DU-DEC-2000; 20-DEC-2000; 20-DEC-2000; 28-PEB-2001; 210-MAY-2001; 210-MAY-2001; 24-PEB-2001; 210-MAY-2001; 24-PEB-2001; 210-MAY-2001; 24-PEB-2001; 210-MAY-2001; 2
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Grimaldi JC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      09
                                                                                                                                                       proteins. The PRO proteins have cytostatic activity. The PRO proteins can be used for targeted delivery of bloactive molecules, such as toxins, radiolabels or antibodies, that cause cell death. PRO nucleotide sequences, and their fragments, can be used as hybridisation probes, in chromosomal and gene mapping, and in the generation of anti-sense RNA
                                                                                                                                                                                                                                      used to develop and screen therapeutically useful reagents. The PRO mucleotide and protein sequence can be used for tissue typing and in treating cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to AAF44470 represent PCR primers and hybridisation probes used in the isolation of human PRO sequences. AAF44087 to AAF4269 and AAB65154 to AAB65300 represent human PRO polynucleotide and protein
                                                                                                                                                                                                                                                                                                                                                                                                                              09
                                                                                                                                                                                                                        and DNA. They may also be used to produce transgenic animals which are used to develop and screen therapeutically useful reagents. The PRO
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                                                                                                                                              transmembrane PRO
                                                               PRO polynucleotides used to produce polypeptides used to target
bioactive molecules such as toxins, radiolabels or antibodies, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; secreted protein; PRO; tumour; lung cancer; colon cancer; breast cancer; prostate tumour; liver tumour; pericyte cell proliferation; chondrocyte cell proliferation; tumour necrosis factor-alpha.
                                                                                                                                                                                                                                                                                                                   sequences given in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 1409; DB 22; Length 265; 100.0%; Pred. No. 2.9e-131; Live 0; Mismatches 0; Indels 0;
                                                                                                                                            present invention describes human secreted and
                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                          specific cells, to cause targeted cell death -
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                                                                                                                 Claim 12; Fig 151; 935pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                        WPI; 2001-032160/04.
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 265; Conserv
                                                                                                                                                                                                                                                                                                                                               265 AA;
                                     N-PSDB; AAF44176
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Godowski PJ; مع Mood WI;

Goddard A,

Watanabe CK,

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The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polynucleotides encode human secreted proteins. The PRO nucleic acids, polypeptides.

CE encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in perioyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, the proliferation or differentiation of chondrocyte cells, for stimulating crimilating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping. AAU83592-AAU83713 represent human PRO protein sequences of the invention.
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                                                                                                         colon cancer, breast tumour, prostate tumour, rectal
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One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours
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Novel isolated PRO polypeptides e.g., PRO826, PRO1068, PRO1184, PRO1346
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Berraran MF, Rong S, Gerber H, Gerriteen ME, Goddard A,
Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J,
Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
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N-PSDB; ABX80266.
11.AVG-2000, 2
23-AVG-2000, 2
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10-JUN-2001, 2
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29-JUN-2001, 2
09-JUL-2001, 2
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25-FEB-1998;
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07-MAY-1998;
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121 IMSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA 180
                                                 NFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLYRSTR 240
                                                                                                                                                                                                                                                                                                 Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing; cardiac insufficiency disorder; cancer; tumour; immune response; adrenal cortical capillary endothelial growth; c-fos induction; vascular endothelial growth factor inhibition; VEGF inhibition; endothelial cell growth inhibitor; T-lymphocytes stimulation; retinal neurons cell survival; rod photoreceptor cell survival; retinal disorder; retinitis pigmentosum; kidney disorder; mammalian kidney mesangial cell proliferation; Berger disease; chondrocyte redifferentiation; sports injury; arthritis.
                            NFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLYRSTR
                                                                                                                                                                                                                                                                           Novel human secreted or transmembrane protein PRO809
                                                                                    RLSEEFGGFRIGNGEVRGRKAAAM 265
                                                                                                    ABUS9107 standard; Protein; 265 AA.
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98WO-US19330.
98WO-US19437.
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2000WO-US03565.
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99WO-US28313,
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16-DBC-1999;
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11-FEB-2000;
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16-SEP-1998;
17-SEP-1998;
07-OCT-1998;
01-DEC-1999;
08-MAR-1999;
02-UUN-1999;
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Godowski PJ; Paoni NF; Wood WI;

DI:

Eaton

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The invention describes an isolated human PRO polypeptide. The PRO polypeptides are useful in detecting PRO polypeptides in a sample, in colypeptides aboutive molecule to a cell expressing a PRO polypeptide, and in modulating at least one biological activity of a cell expressing a PRO polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus useful for treating cardiac insufficiency disorders. PRO154 and PRO1186 ct. PRO943, PRO828, PRO826, PRO943, PRO828, PRO826, PRO943, PRO826, PRO943, PRO828, PRO826, PRO943, PRO826, PRO943, PRO928, PRO943, PRO944, PR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and therefore are useful for treating kidney disorders associated with decreased mesangial cell function such as Berger disease or other nephropathies associated with dermatitis, herpetiformis or Crohn's disease. PRO1310, PRO844, PRO1312, PRO1192 and PRO1387 induce the proliferation and/or redifferentiation of chondrocytes in culture and are thus useful for treating sports injuries, and arthritis. This is the amino acid sequence of a novel human PRO protein.
and PRO1375, which stimulate proliferation of stimulated T-lymphocytes are therapeutically useful for enhancing immune response and in cancer
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                                                                                                              Claim 12; Fig 151; 648pp; English
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Best Local Similarity
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Human; PRO; secreted; transmembrane; pharmaceutical; diagnostic; blosensor; bloreactor; tumour; therapeutic, gene therapy; tumour-associated antigenic target; TAT; ADEPT; antibody-dependent enzyme mediated prodrug therapy; cytostatic.
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97US-065311P.
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99WO-US21547.
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Matches 265; Conserv
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Grimaldi JC,
              13-NOV-1997;
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25-FEB-1998;
26-MAR-1998;
77-MAY-1998;
02-JUN-1998;
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Roy MA, ...
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Godowski PJ; Paoni NF; Wood WI; J, Baker KP, Botstein D, Desnoyers L, Eaton J Fong S, Gerber H, Gerritsen NE, Goddard A, Gurney AL, Kljavin IJ, Napier MA, Pan J, J ewart TA, Tumas D, Watanabe CK, Williams PM,

New transmembrane polypeptides and nucleic acids encoding the polypeptides, useful in gene therapy, in chromosome identification, as chromosome markers, or in generating probes -

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The invention discloses isolated PRO secreted/transmembrane polypeptides comprising a sequence without signal peptide and the nucleic acid mencéding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. The PRO polypeptides or
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polynucleotides are also useful in gene therapy, in chromosome lidentification, as chromosome markers, or in generating probes. The PRO polypeptides are useful as molecular markers for protein electrophoresis, and the isolated nucleic acids may be used for recombinantly expressing those markers. The PRO polypeptides and nucleic acids may also be used in tissue tryping. Anti-PRO antibodies are useful in diagnostic assays for PRO, and in affinity purification of PRO from ABUG0478-ABUG0624 are the PRO polymocleotides of the invention.

Note: The sequence data for this patent is also available in electronic format from USPTO at seqdata.uspto.gov/sequence.html.
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horse; cow; dog; cat; sheep; pig; goat; rabbit; ADEPT;
antibody-dependent enzyme mediated prodrug therapy.
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100.0%; Score 1409; DB 24; Length 265;
Best Local Similarity 100.0%; Pred. No. 2.9e-131;
Matches 265; Conservative 0; Mismatches 0; Indels 0;
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98WO-US19437.
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PR 16-DEC-1999) 99W0-US28634
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PR 24-MAC-2000 2000W0-US0341
PR 24-MAC-2001 2000W0-US0341
PR 24
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98US - 0855 8 P.
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17-AUG-1998; 18-AUG-1998; 18-AUG-1998;

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08-AAN-1999; 99WO-US25108.
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02-JUN-1999; 99WO-US26028.
01-DEC-1999; 99WO-US21907.
01-DEC-1999; 99WO-US28313.
02-JAN-2000; 2000WO-US30211.
02-JAN-2000; 2000WO-US03414.
02-MAR-2000; 2000WO-US03414.
02-MAR-2000; 2000WO-US03414.
02-MAR-2000; 2000WO-US03414.
02-MAR-2000; 2000WO-US03413.
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03-MAR-2000; 2000WO-US03413.
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03-MAR-2000; 2000WO-US132678.
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pharmaceutical, diagnostic, biosensor, bioreactor, tumour, therapeutic, colon cancer, lung cancer, breast cancer, gene therapy.
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al Similarity 100.0%;
265; Conservative 0
98US-096959P.
98US-096960P.
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98US-097161P.
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Best Local Similarity
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17-SEP-1998;
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Homo sapiens.
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28-JUL-2000;
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                                                                                                                                                            ABU13920;
                                                                                        RESULT 11
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                                                                                                                                           The invention discloses isolated PRO secreted/transmembrane polypeptides comprising a sequence without signal peptide and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. The PRO polypeptides or polymucleotides are also useful as pharmaceuticals, diagnostics, biosensors or bioreactors, for detecting or treating e.g. tumours in mammals, e.g. humans, dogs, cats, cattle, horses, sheep, pigs, goats or rabbits as targets for therapeutic intervention in certain cancers (e.g. colon, lung or breast cancers) and diagnostic determination of the presence of these cancers. The PRO polypeptides are also useful as molecular weight markers or for chromosome identification. The PRO genes are useful as hybridisation probes or for servening libraries of human cDNA, genomic DNA, or mRNA. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequences presented in ABUSB900-ABUS9046 are the PRO polypeptides of the invention.
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                                                                                                                                                                                                                                                                                                                                            Godowski PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New secreted and transmembrane PRO polypeptides (e.g. PRO183, PRO184, PRO361 or PRO846) useful as targets for therapeutic intervention in cancers (e.g. lung or breast cancers), or for diagnosing these cancers
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Wood WI;
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Ferrara N, Fong S, Gerber H, Gerriteen ME, Goddard A,
Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J,
Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
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                                98US-089440P.
98US-089512P.
98US-089514P.
98US-089532P.
98US-089538P.
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98US-089600P.
98US-089653P.
98US-088876P.
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17-JUN-1998;
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16-JUN-1998
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Human; PRO polypeptide; secreted protein; transmembrane protein;
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241 RLSEEBFGGFRIGNGEVRGRKAAAM 265
                        241 RLSEEEFGGFRIGNGEVRGRKAAAM
                                                                                                                                 ABU13920 standard; Protein; 265
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98WO-US19437
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                                                                                                                                                                                                              26-FEB-2003 (first entry)
                                                                                                                                                                                                                                                   Human PRO809 polypeptide.
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97US-065311P.
97US-065710P.
97US-06570P.
98US-078910P.
98US-08322P.
98US-081106P.
98US-087106P.
98US-08750P.
98US-08750P.
98US-08752P.
98US-08752P.
98US-08752P.
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                        02-JUN-1998;
02-JUN-1998;
02-JUN-1998;
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16-JUN-1998;
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16-JUN-1998;
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04-JUN-1998
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18-JUN-1998
   NOV-1997
                                                                           05-JUN-1998
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(GETH) GENENTECH LTD.

Godowski PJ; Paoni NF; Wood WI; Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL, Ferrara N, Rong S, Gerber H, Gerritsen ME, Goddard A, God Grimaldi JC, Gurney AL, Kljavin IJ, Napire MA, Pan J, Paor Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Woo 7 Zhang

WPI; 2003-102117/09.

N-PSDB; ABX64090.

Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers

Claim 12; Fig 151; 649pp; English.

The present invention relates to the isolation of novel human PRO polypeptides, and the polymucleotide sequences encoding them. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for for identifying agonists or antagonists. The polymucleotide sequences encoding PRO polypeptides are useful as

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; PRC; secreted polypeptide; transmembrane polypeptide;
toxin; radiolabel; cell death; gene mapping; chromosome mapping;
protein electrophoresis; genetic disorder; immunosuppressive; cytostatic;
antibacterial.
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hybridisation probes, in chromosome and gene mapping, in the generation of antisense RNA and DNA, in the preparation of PRO polypeptides, for generating transgenic animals or knockout animals, to construct hybridisation probes for mapping the gene which encodes the PRO polypeptide, and for the genetic analysis of individuals with genetic disorders, in gene therapy, for chromosome identification, as chromosome markers, and for generating probes for PR. Northern analysis, Southern analysis and Western analysis. ABU13860-ABU14006 represent the human PRO polypeptides of the invention.

Note: The sequence data for this patent was obtained in electronic format directly from the USPTO web site at sequence analysis substitution.
                                                                                                                                                                                                                                                                                                                                                                                                 61 SLCGTKNIKVAKKVVKTHEPASFNINVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE
                                                                                                                                                                                                                                                                                                                                 1 MGLEGLFCLAVLAASSFSKAREEBITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPFITY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121 LWSKPVSELRANFTLØDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA
                                                                                                                                                                                                                                                                                                             1 MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPPITY
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                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                             100.0%; Score 1409; DB 24; Length 265; 100.0%; Pred. No. 2.9e-131;
                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     241 RISEEFFGGFRIGNGEVRGRKAAAM 265
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98WO-US19437
98WO-US21141
99WO-US00106
99WO-US0028
99WO-US05028
99WO-US05028
99WO-US1090
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                                                                                                                                                                                                               265 AA;
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17-SEP-1998;
07-OCT-1998;
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02-JUN-1999;
15-SEP-1999;
15-SEP-1999;
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2000WO-US05841.
2000WO-US06319.
2000WO-US06884.
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98US-084600P.
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98US-087607P.
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9805-088021P.
9805-088025P.
9805-088025P.
9805-088026P.
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98US-088033P.
98US-088132EP.
98US-088167P.
98US-088212P.
98US-088212P.
98US-08855SP.
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97US-066770P.
98US-075945P.
98US-078910P.
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98US-088824P.
98US-088826P.
98US-088856P.
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98US-089512P.
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2000WO-US13358.
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2001WO-US06520.
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2001WO-US21735.
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97US-062250P
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98US-089105P.
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99WO-US28301
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30-NOV-1999;
01-DBC-1999;
16-DBC-1999;
20-DBC-1999;
20-DBC-1999;
20-DBC-1999;
20-DBC-1999;
20-DBC-1999;
21-FEB-2000;
22-FEB-2000;
22-FEB-2000;
24-FEB-2000;
24-FEB-2000;
26-MAR-2000;
20-MAR-2000;
20-MAR-2000;
20-MAR-2000;
20-MAR-2000;
21-MAY-2000;
21-MAY-2000;
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27-MAY-2000;
28-MAY-2000;
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20-MAY-2000;
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09-JUL-2001;
16-JUN-1997;
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25-FEB-1998;
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07-MAY-1998,
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The invention relates to a secreted and transmembrane polypeptide, termed FRO polypeptide, and the polynucleotide encoding it. The polypeptide is useful for detecting PRO polypeptides and for linking a bioactive molecule to a cell expressing the above polypeptides, where the bioactive molecule is a toxin, radiolabel or an antibody. The bioactive material causes the death of the cell. The polypeptide is useful for identifying agonists or antagonists of the PRO polypeptide, for preparing variants of PRO, as a molecular weight marker for protein electrophoresis purposes and the PRO polynucleotide is useful for recombinantly expressing those markers. The polynucleotide is also useful as a hybridisation probe, in chromosome and gene mapping, in generation of antisense RNA and DNA, in the preparation of PRO polypeptide, for generating transgenic animals or thorowing in the mapping, in generation of antisense RNA and DNA, in the preparation of therapeutically useful reagents, to construct hybridisation probes for mapping the gene which encodes PRO and for the genetic consumpting the gene which encodes PRO and for the genetic consumptions of individuals with genetic disorders, in gene therapy, for chromosome identification, as a chromosome marker and for generating probes for PRO, Northern analysis, Southern analysis and Western analysis. This sequence represents a human PRO polypeptide of the
                                                                                                                                                                                                                                                             DL;
Godowski PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers -
                                                                                                                                                                                                                                                                                                 Paoni NF;
Wood WI;
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Bernara N, Fong S, Gerber H, Gerriteen ME, Goddard A,
Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J,
Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM,
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                98US-089538P.
98US-089598P.
98US-089509P.
98US-089600P.
98US-089653P.
98US-089601P.
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28-AUG-2001; 2001US-0941992
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                                     17-JUN-1998;
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                                                                                                                     .8-JUN-1998;
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SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE 120 SLCGTKNIKVAKKVVKTHEPASFNIMVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE 120 9 09 1 MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPFITY MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPFITY 0; Gaps 24; Length 265; 0; Indels 100.0%; Score 1409; DB 24; 100.0%; Pred. No. 2.9e-131; tive 0; Mismatches 0; Query Match 100. Best Local Similarity 100. Matches 265; Conservative 61 a à

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LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA 180

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121 LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA

181 NESFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKWEDWQGPLESPILALPLYRSTR 181 NFSFLPSGTSDWFWCQAANMANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLYRSTR

RISEEFFGGFRIGNGEVRGRKAAAM 265

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EST encoded protein SEQ ID NO: 1997.
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                                                                                                                                                       AAM24472 standard; Protein; 232 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JAN-2000; 2000US-0491404.
17-JUL-2000; 2000US-0617746.
03-AUG-2000; 2000US-0631451.
15-SEP-2000; 2000US-063870.
                                                                                                                                                                                                                                                                                                                                                                                                                             25-JAN-2001; 2001WO-US02687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibodies and research use
                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 64.3
Matches 148; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cao Y, Drmanac RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-476164/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           232 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAH99131
                                                                                                                                                                                                                                                                                                                                                                     WO200154477-A2
                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                              12-0CT-2001
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                                                                                                                                                                                    AAM24472;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to full length GENSET human nucleic acids encoding potentially secreted proteins. The nucleic acids and the polypeptides they encode may be used in the prevention, treatment and diagnosis of example, they be used in the prevention, treatment and diagnosis of example, they be used to treat disorders associated with decreased GENSET gene expression. For example, they be used to treat disorders associated with decreased GENSET gene expression by rectifying mutations or deletions in a patient's genome that affect the activity of GENSET or by supplementing the patients own production of GENSET polypeptides. Conversely, antisense nucleic acid molecules may be administered to down regulate GENSET expression. The sense and antisense nucleic acid sequences in samples, and hence to be presence of similar nucleic acid sequences in samples, and hence to determine which patients may be used as antigens in the production of antibodies and in assays to identify modulacrs (agonists and in any modulacrs (agonists and in any modulacrs (agonists and in any modulacrs).
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                                                                                                                                                                                 secreted protein; gene therapy; vaccine; treatment; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Full length GENSET human nucleic acids encoding potentially secreted proteins, useful in gene therapy and vaccination against a variety of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antagonists) of GENSET polypeptide expression and activity. The present sequence is a GENSET polypeptide of the invention.
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100.0%; Pred. No. 1.6e-105;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins, useful in gene therapy and vaccinatior
diseases, and for diagnosis of those diseases -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 21; Page 827-828; 921pp; English.
                                                                                                                                                      Human secreted protein, SEQ ID NO: 296
241 RLSEEFFGGFRIGNGEVRGRKAAAM 265
                                                                      ΑA
                                                                  AAG89176 standard; Protein; 247
                                                                                                                                                                                                                                                                                                             07-DEC-2000; 2000WO-IB01938.
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Best Local Similarity
Matches 215; Conserv
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                                                                                              AAG89176;
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                                                                                                                                                                                                 GENSET
                                       RESULT 13
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121 LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse; tomato; monkey; dog; sea urchin; expressed sequence tag; EST; diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Asundi V;
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                                                                                                                                          181 NFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGG 215
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Zhang J, Werhman T;
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LWSRQRG-----RPQGGDDLPGVLGQPTYHQQPDREGWAGPPAAETMPQEACQLSPS 172
LWSKPVSELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention comprises the amino acid and coding sequences of human secreted proteins. The DNA and protein sequences of the invention are useful for the diagnosis and treatment of allergic disorders, asthmatic disorders and immediate hypersensitivity diseases (e.g. hay fever, allergic conjunctivitis and allergic rhinitis). The proteins of the invention are also useful for identifying a binding partner. The nucleic acids of the invention are also useful for chromosome identification, radiation hybrid mapping or long-range restriction mapping. The present amino acid sequence represents a human secreted protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of human secreted proteins and nucleic acids for preparing a diagnostic or pharmaceutical composition for diagnosing or treating allergic or asthmatic disorders, e.g. asthma, hay fever, or allergic conjunctivitis or rhinits -
                                                                                                                                                                                                                                                                                                                                                       Human; protein therapy; immediate hypersensitivity disease; allergic disorder; asthmatic disorder; gene therapy; secreted hay fever; allergic conjunctivitis; allergic thinitis; binding partner identification; chromosome identification;
                                                                                                           ----CRARHRIWFWCQACKQRQCSSTAPSQWLPQVVTQKMEDWQGPPGEP
                                                                          181 NFSFLPSQTSDWFWCQAANNANVQHSALTV-VPPGGDQKMEDWQGPLESP
                                                                                                                                                                                                                                                                                                                      Human secreted protein amino acid sequence - SEQ ID No 148.
                                                                                                                                                                                                                                                                                                                                                                                                                                 radiation hybrid mapping; long-range restriction mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 632-633; 823pp; English.
                                                                                                                                                                                                        ABJ19682 standard; Protein; 235 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-2001; 2001US-278650P.
12-SEP-2001; 2001US-0950082.
12-SEP-2001; 2001US-0950083.
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                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                              binding partner
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 121
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1 MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPITY 1 MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPITY

46.5%; Score 654.5; DB 24; Length 235; 59.3%; Pred. No. 1.8e-56; live 10; Mismatches 44; Indels 47;

Conservative

Local Similarity Les 147; Conserv

Best Loca Matches

Query Match

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The invention relates to novel human genes (ABZ66891-ABZ68209) and the encoded secreted proteins (ABP99470-ABP99872) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and in the specification. The nucleic acids, proteins, antibodies and in the specification. The nucleic acids, proteins, antibodies and antibacing are useful in the diagnosis, treatment and prevention of:

(a) cancer, e.g. breast and ovarian cancer and other cancers of the arenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, ung or urogenital; (b) immune disorders e.g. Addison's disease,

(a) lergies, autoimmune haemolytic anneand, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid architis; (c) cardiovascular disorders such as viral, cerebral anoxia and epilepsy; and (f) infectious diseases such as viral,
                                                                                                                   200
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; secreted protein; nootropic; neuroprotective; cytostatic; viruide; dermatological; immunosuppressive; antiinflammatory; anti-HIV; vulnerary; antibacterial; antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antirheumatic; hepetotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; infection; vaccine; cardiovascular disorder; neurological disease; nephrotropic;
121 LWSKPVSELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGOVHLQORPCHR--- 177
                                   -QRGRPQGGDDLPGVLGQPTYHQQPDREGWA
                                                                             178 -QPANFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLY
                                                                                                    New human secreted proteins encoded by genes contained in cDNA clones (e.g. HGCAC19), useful for preventing, treating or diagnosing e.g. ALIDS, multiple sclerosis, herpes virus, leukemia, tick-boxne encephalitis or West Nile fever
                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted protein SEQ ID NO 516.
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                                                                                                                                                                                                                                                                                                ABP99572 standard; Protein; 235
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12-SEP-2001; 2001US-0950082.
12-SEP-2001; 2001US-0950083.
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N-PSDB; ABZ66993.
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                                     121 LWSR-
                                                                                                                   155 GPPA
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RESULT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; secreted protein; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; oytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; neoplasm; opthalmalogical; autoimmune disease; rheumatoid arthritis; anglogenesis; opthalmalogical; autoimmune disease; rheumatoid arthritis; anglogenesis; cerebrovascular disorder; cardiovascular disorder; nervous system disorder; infection; wound healing; chemotaxis.
                                                                                                                                                                                                                       177
                                                                                                                                                              SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE 120
                                                                                                                                                                                        61 SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE 120
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                                                                                                         1 MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPITY
                                                                                                                                   MGLPGLFCLAVLAASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPPITY
                                                                                                                                                                                                                    LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHR---
                                                                                                                                                                                                                                               --- QRGRPQGGDDLPGVLGQPTYHQQPDREGWA
                                                                                                                                                                                                                                                                          -QPANFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLY
                                                                                                                                                                                                                                                                                                    ----AETMPQAACQLLLPAEPDIGLVLV--PGCKQR----QCPAQRPHSGAP--
                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding a human secreted protein is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted protein sequence encoded by gene 38 SEQ ID NO:96.
                                                                                  47;
                                                       DB 24; Length 235;
                                                                                 Indels
                                                                                 44;
                                                      Score 654.5; DB 24
Pred. No. 1.8e-56;
                                                46.5%; Score 03...,
59.3%; Pred. No. 1.8e-
+ive 10; Mismatches
 parasitic infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 11; Page 386-387; 434pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Komatsoulis G;
                                                                                                                                                                                                                                                                                                                                                                                                                            AAB39216 standard; Protein; 236 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L6-MAR-2000; 2000WO-US06792.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0125362.
99US-0169980.
                                                                 Local Similarity 59.3%
les 147; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                               121 LWSR-----
bacterial, fungal and
                                                                                                                                                                                                                                                                                                                              237 RSTRRLSE 244
                                                                                                                                                                                                                                                                                                                                                          ---RRVXQ 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rosen GA, Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-579483/54
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                                                                                                                                                                                                                                                                                                    155 GPPA-
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                            Sequence
                                                       Query Match
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AAB39216
                                                                                 Matches
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protection based on the tissues and cells in which they are expressed.

Examples of activities include: immunosuppressive; antiathritic;

Examples of activities include: immunosuppressive; antiathritic;

Examples of activities include: immunosuppressive; antiathritic;

carebroprotective; nootropic; neuroprotective; antibacterial; virucide;

fungicide; and opthalmalogical. The human secreted proteins,

continued, prevention, and/or diagnosis of the invention may be useful

in the treatment, prevention, and/or diagnosis of various disease.

Cliver, populations such as autoimmune diseases e.g. rheumatoid

atthritis, hyperproliferative disorders e.g. neoplasms of the breast or

liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular

disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders

c.g. Altelmer's disease, infections caused by bacteria, viruses and

c.g. Altelmer's disease, infections caused by bacterial, viruses and

c.g. also be used to aid wound healing and epithelial cell proliferation, to

cegenerate tissues, maintain organs before transplantation, in

cementaria and as a food additive or preservative e.g. to increase

chemotaxia and as a food additive or preservative e.g. to increase

contract contraction and characterisation of the genes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        178 -QPANFSFLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 SLCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LWSKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHR---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ---QRGRPQGGDDLPGVLGQPTYHQQPDREGWA
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   The proteins have
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Alzheimer's disease, infection, ocular disorder, corneal infection;
wound healing; epithelial cell proliferation; food additive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hyperproliferative disorder; breast neoplasm; cancer; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; foetal tissue antigen; antiinflammatory; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 46.5%; Score 654.5; DB 21; Length 236;
59.3%; Pred. No. 1.8e-56;
iive 10; Mismatches 44; Indels 47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44; Indels
protein sequences with which they share homology.
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---RRVXQ 205
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|21 LWSR-
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PR 02-0CT-2000; 2000US-0234037.
PR 02-0CT-2000; 2000US-0237037.
PR 02-0CT-2000; 2000US-0237039.
PR 13-0CT-2000; 2000US-0237039.
PR 13-0CT-2000; 2000US-0239337.
PR 20-0CT-2000; 2000US-0239337.
PR 20-0CT-2000; 2000US-0239337.
PR 20-0CT-2000; 2000US-0241785.
PR 20-0CT-2000; 2000US-0241785.
PR 20-0CT-2000; 2000US-0241786.
PR 20-0CT-2000; 2000US-0241787.
PR 20-0CT-2000; 2000US-0241787.
PR 20-0CT-2000; 2000US-0241826.
PR 20-0CT-2000; 2000US-0246474.
PR 20-0CT-2000; 2000US-024622.
PR 20-0CT-2000; 2000US-024221.
PR 20-0
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New polynucleotides and polypeptides for diagnosing, treating,

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The invention relates to novel nucleic acids encoding novel human foetal antigens. The nucleic acids and proteins are used to prevent, treat (e.g. by gene therapy) or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They care also used in diagnosing a pathological condition or susceptibility or a pathological condition. The antibodies to the antigens can also be used in alleviating symptoma associated with the disorders and in diagnostic immunoassays e.g. radioimmunossays or enzyme linked immunoassays (ELISA). Disorders which are diagnosed or treated infunde autoinmune diseases e.g. rheumatoid arthritis, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sumburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also cas a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, minerals, cofactors and other nutritional components. Mumerous examples of diseases and disorders treated by the nucleic acids and proteins are given in the specification. The present sequence
        the nervous,
                                excretory, gastrointestinal, reproductive, and
     e.g. diseases or disorders of
                                                                                                         Claim 11; SEQ ID No 1500; 642pp; English
     or prognosing
                                                          respiratory systems
                             musculoskeletal,
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                                                                                                                                 107
                                                                                                                                                                                                               108 HEPASFINIAVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWELWSKPVSELRANFTLOD 167
                                                                                         77
                                                                                                                                                                    78 HEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSKPVSELRANFTLQD
                                                                                    18 SKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPPITYSLCGTKNIKVAKKVVKT
                                                                                                                            48 SQSPSLEITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPPITYSLCGTKNIKVAKKVVKT
                                              0; Gaps
45.1%; Score 636; DB 22; Length 175; 96.0%; Pred. No. 8.2e-55; ive 2; Mismatches 3; Indels (
                    Best Local Similarity 96.0 Matches 120; Conservative
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Human immunoglobulin receptor isoform IRTA2a.
                                                                                                                                                 AAB82313 standard; Protein; 759 AA.
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RESULT 19
AAB82313
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Immunoglobulin superfamily receptor translocation associated; IRTA; IRTA2a; human; immunoglobulin receptor; Fc receptor; melanoma; lymphoma; myeloma; B cell malignancy; cancer; chromosome 1q21; diagnosis; therapy. Homo sapiens.

132.,134 /note= "Asn is N-glycosylated" 383.,385 ...15 'label= Signal_peptide 16..759 // label= Mature_protein Location/Qualifiers Peptide Protein

Modified-site

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The present sequence is that of the novel human immunoglobulin receptor, immunoglobulin superfamily receptor translocation associated protein isoform 2a (IRTA2a), an Fc receptor involved in carbon sequence is electronically secondated protein isoform 2a (IRTA2a), an Fc receptor involved in the pathogenesis of lymphoma and melanoma. Efforts to identify compared in chromosomal aberrations affecting band [q21] in multiple myeloma and B cell lymphoma led to the discovery of IRTA2 and IRTA1 (see AAB82312) as founding members of a novel subfamily conforms in transcribed into 3 major mRNA isoforms, IRTA2a is a 759 amino conforms in transcribed into 3 major mRNA isoforms, IRTA2a is a 759 amino conforms in the B cell supplementation of a unique C-terminus. IRTA2b diverges from IRTA2a at residue 560, extending for a further 31 residues. The IRTA2 diverges from IRTA2a at cesting 560, extending for a further 21 residues. The IRTA2 genes CC residue 746 and extends for a further 21 residues. The IRTA2 genes CC residue 746 and extends for a further 21 residues cells, which may include immunoblasts and memory cells. The invention provides IRTA2 is expressed in GC centrocytes and in perifolicular cells, which may conclude immunoblasts and memory cells. The invention provides IRTA2 include immunoblasts and memory cells. The invention provides IRTA2 conclude immunoblasts and memory cells. The invention provides IRTA2 conclude immunoblasts and memory cells. The invention provides IRTA3 conclude immunoblasts and memory cells. The invention provides IRTA3 conclude immunoblasts and memory cells. The invention provides IRTA3 conclude immunoblasts and anticitally hybridises with a unique coll conclude that specifically hybridises to IRTA mEMA so as to prevent oversexpression of IRTA protein and hence to arrest coll growth or induce cell drom B cell lymphoma, multiple myeloma, Burkitt's lymphoma, martiple myeloma, Burkitt's lymphoma, martiple myeloma, Burkitt's lymphoma, martiple myeloma, Burkitt's lymphoma, martiple myeloma, Burkitt's 64 GTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGA-HVDSARLQMHWELW 122 13 AASSFSKAREEEITPVVSIAYK----VLEVFPKGRWVL----ITCCAPQPPPITYSLC New genes encoding immunoglobulin receptor, Immunoglobulin super Receptor Translocation Associated proteins, used to treat B cell malignancies including lymphomas and multiple myeloma -DB 22; Length 759; Indels 40; Mismatches 105; /note= "Asn is N-glycosylated" 621..623 /note= "Asn is N-glycosylated" /note= "Asn is N-glycosylated" 714.716 /note= "Asn is N-glycosylated" 8.1%; Score 114.5; DB 25.1%; Pred. No. 0.038; Claim 3; Fig 18B-1-18B-2; 72pp; English. (UYCO) UNIV COLUMBIA NEW YORK, 99US-0168151 28-NOV-2000; 2000WO-US32403 Local Similarity 25.1 631..633 WPI; 2001-355921/37. 759 AA; N-PSDB; AAF30950 Jalla-Favera R; WO200138490-A2 Modified-site Modified-site Modified-site 29-NOV-1999; 31-MAY-2001. Sequence Query Match Best Loca Matches g ð à

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                SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
602 -HEDVTLGSSSAPSGGEASFNLSLTAEHSGN---YSCEANNGLVAQHSDTISLSVIVPV- 656
                              657 SRPILTFRA--PRAQAVVGDLLELHĆEALRĠSSPILYWFYHEDVTLGKISAP-SGGGASF
                                                                                                                                                          Immunoglobulin superfamily receptor translocation associated; IRTA;
IRTA2c; human; immunoglobulin receptor; Rc receptor; melanoma;
lymphoma; myeloma; B cell malignancy; cancer; chromosome 1q21;
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                                             183 SF-LPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLES 228
                                                      /note= "putative consensus Src-homology
binding domain"
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te= "Asn is N-glycosylated"
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                                                                                                                                                                                                                                                              "transmembrane domain"
                                                                                                                                            Human immunoglobulin receptor isoform IRTA2c.
                                                                                                                                                                                                                               /label= Signal_peptide
16..977
/label= Mature_protein
                                                                                                                                                                                                                                                                                                                                                                                                                          binding domain"
                                                                                                                                                                                                                 iocation/Qualifiers
                                                                                                AAB82315 standard; Protein; 977 AA
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806..808
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899..902
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                                                                                                                                                                                                                                                                                     383
                                                                                                                                                                                    diagnosis; therapy
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                123
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Peptide
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The present sequence is that or the novel numminoglobulin superfamily receptor translocation associated protein isoform 2c (IRTRAC), an PC receptor involved in the pathogenesis of lymphoma and melanoma. Efforts to identify genes involved in chromosomal aberrations affecting band 1q21 in multiple myeloma and B cell lymphoma led to the discovery of IRTRAC of the IRTRAC is a founding members of a novel subfamily of related receptors within the immunoreceptor family. The IRTRAC of related receptors within the immunoreceptor family. The IRTRAC of IRTRAC (see also AAB82312) and AAB82314). IRTRAC is the longest is transcribed into 3 major mRNA isoforms, IRTRAC and IRTRAC (see also AAB82313) and AAB82314). IRTRAC is the Industry is a type I transmembrane glycoprotein. Each SH2 binding site agrees with the immune receptor tyrosine-based inhibition motif (ITIM) consensus and is encoded by a separate exon. The IRTRAC is expressed in GC centrocytes and in perifolaliquar cells, which may include immunoblasts and memory cells. The invention provides IRTA nucleic acids and proteins, and antibodies directed to epitopes of IRTA proteins Methods are claimed for: detecting a B cell malignancy comprising a 1q21 chromosomal rearrangement using a collone of human IRTAL-5; and treating a subject having a B cell cancer by administering an anti-IRTA antibody or an antisense collower of human IRTAL-5; and treating a subject having a B cell concer by administering an anti-IRTA antibody or an antisense collower or induce cell death of cancer cells expressing IRTA. The lamboms and matched and marking and have an antisense collower. Mantle and antition may mark and manthems and manthems and manthems and marking and having an anti-IRTA protein and hence to arrest to prevent overexpression of IRTA protein and encer cells expressing IRTA. The lamboms marking marking any anti-IRTA lamboms marking and any anti-IRTA lamboms marking and any and 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      602 -HEDVTLGSSSAPSGGEASFNLSLTAEHSGN---YSCEANNGLVAQHSDTISLSVIVPV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  657 SRPILTFRA--PRAQAVVGDLLELHCEALRGSSPILYWFYHEDVTLGKISAP-SGGGASF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64 GTKNIKVAKKVVKTHEPASFNINVTLKSSPDLLTYFCRASSTSGA-HVDSARLQMHWELW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF
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                                                                                                                                                                                            New genes encoding immunoglobulin receptor, Immunoglobulin super
Receptor Translocation Associated proteins, used to treat B cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 the novel human immunoglobulin
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                                                                                                                                                                                                                                           Receptor Translocation Associated proteins, used to tr
malignancies including lymphomas and multiple myeloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.8%; Score 110.5; DB 22; 24.6%; Pred. No. 0.13; ive 41; Mismatches 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : | :: | : | : | : | : | : | : | 14 NLSLTTEHSGIYSCEADNGLEAQRSEMVILKVA----
                                                                                                                                                                                                                                                                                                                                                           Claim 3; Fig 18B-1-18B-2; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                 WPI; 2001-355921/37
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                                                                                                                            N-PSDB; AAF30952
Dalla-Favera R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present
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Matches
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marrow; umbilical cord venous endothelial cell;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention provides polypeptides (AAY27127-Y27133) produced by human adult brain tissue, human bone marrow or a human umbilical cord venous endothelial cell. Host cells transformed with vectors comprising the mucleic acids encoding the polypeptides are used for the recombinant diagnosis, treatment and basic studies, with wide applications in treatment depending on the activity to be aimed at. Sequences AAX89112-125 represent nucleic acids encoding the polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89 LKSSPDLLTYFCRASSTSGAH------VDSARLQMHWELWSKPVSELRANFTLQDRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SE HESGP----YKCKAQVISCSKYSRDFSFIIVDPV----SF-ISPVINIMVIQTETDR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  37 EVFPKGRWVLITCCAPQPPPITYSL-----CGTKNIKVAKKVVKTHEPASFNLNVT--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44 KVVMKGQNVSMFCSHKNKSLQITYSLFRRKTHLGTQDGK------GEPALFNLSITEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                               204 QHS-ALTVVPPGGDQKMED-----WQ--GPLESPILALPLYRSTRRLSEE 245
                                                                                                                            377 OCSEAVPVSISGPDGYRRDIMTAGVLWGLFGVLGFTGVALLLYALFHKISGE 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 343;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New adult human brain tissue-produced polypeptides useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        67; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Human bone marrow-derived polypeptide (clone OAF038-Leu).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 7.2%; Score 101.5; DB Best Local Similarity 26.4%; Pred. No. 0.24; Matches 55; Conservative 27; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "signal peptide"
20..343
/note= "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 54-55; 86pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                 AAY27129 standard, Protein, 343 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tada H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recombinant; diagnosis; treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; AAX89116, AAX89117
                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                         tissue; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-419088/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         343 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fukushima D,
                                                                                                                                                                                                                                                                                                                                             14-SEP-1999
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                                                                                                                                                                                                                                                                                                AAY27129;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein
                                                                                                                                                                                                  RESULT 22
                                                                                                                                                                                                                          AAY27129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     145 EMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANFSF-LPSQTSDWFWCQAANNANV 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IRTA genes display a specific pattern of expression in mature B cells. IRTA4 is selectively expressed in mantle zone C cells, the pre-GC compartment of mature B cells. The invention provides IRTA nucleic acids, proteins, and antihodies directed to epitopes of IRTA proteins. Methods are claimed for: detecting a B cell malignancy comprising a lq21 chromosomal rearrangement using a nucleic acid momen in a pacifically hybridises with a unique sequence of human IRTA1-5, and treating a subject having a B cell cancer by administering an anti-IRTA antibody or an antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of the novel human immunoglobulin receptor, immunoglobulin superfamily receptor translocation associated protein 4 (IRTA4), an Fc receptor involved in the pathogenesis of lymphoma and melanoma. Efforts to identify genes involved in chromosomal aberrations affecting band 1g21 in multiple myeloma and B cell lymphoma led to the discovery of IRTA1 and IRTA2 (see ABB2312-15) as founding members of a novel subfamily of related receptors within the immunoreceptor family. 3 Additional proteins, IRTA3, IRTA4 and IRTA5 (see ABB2316-18), were subsequently identified, which are also members of this novel subfamily. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     218 EGOKLILLCSVAGGTGNVTFSWYREATGTSMGK-----KTORSLSAELBIPAVKESDAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 KGRWVLITCCAPQPPPPITYS----LCGTKNIKVAKKVVKTHEPASFNLNVTLKSSPDLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97 TYFCRASSISGAHVDSARLOMHWELWSK-----PVSELRANFILQDRGA----GPRV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   48; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple myeloma, Burkitt's lymphoma, marginal zone lymphoma, diffuse large cell lymphoma and follicular lymphoma. The B cell lymphoma is selected from mucosa-associated-lymphoid tissue B cell lymphoma or non-Hodgkin's lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              that specifically hybridises to IRTA mRNA so as to prevent overexpression of IRTA protein and hence to arrest cell growth or induce cell death of cancer cells expressing IRTA. The B cell cancer is selected from B cell lymphoma, mantle cell lymphoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New genes encoding immunoglobulin receptor, Immunoglobulin super
Receptor Translocation Associated proteins, used to treat B cell
malignancies including lymphomas and multiple myeloma -
                                               "mwunoglobulin superfamily receptor translocation associated,
"RTA4; human; immunoglobulin receptor; Fc receptor; melanoma;
ymphoma; myeloma; B cell malignancy; cancer; chromosome 1q21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 22; Length 508;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              :96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunoglobulin receptor IRTA4 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5; Fig 18D-1-18D-2; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                          28-NOV-2000; 2000WO-US32403
                                       Immunoglobulin superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001-355921/37.
                                                                                                                    diagnosis; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      272 KYYCRADNG-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  508 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAF30954
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dalla-Favera R;
                                                                                                                                                                                                              WO200138490-A2
                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                         29-NOV-1999;
                                                                                                                                                                                                                                                             31-MAY-2001
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Matches
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13;

88

à qq à g

g à g

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57.59 -
/hoce= "protein kinase C phosphorylation site"
71.74 -
/note= "cAMP- and cGMP-dependent protein kinase
                                                                                                                                                                                                                                                        TANGO 228; human; spleen disorder; immunological disorder; immunomodulator; antiinflammatory; cancer; tumour; metastasis; antitumour; anticancer; antimetastatic; therapy; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note= "protein kinase C phosphorylation site"
84..186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "protein kinase C phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "protein kinase C phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note= "protein kinase C phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91. 94
/note= "casein kinase phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "casein kinase phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "protein kinase C phosphorylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Asn is N-glycosylated"
157..160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51..54
/note= "Asn is N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60..63
/note= "Asn is N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'note= "Asn is N-glycosylated"
82..185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Asn is N-glycosylated"
18..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "immunoglobulin domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Asn is N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'label= immunoglobulin domain
                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "extracellular domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           "transmembrane domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorylation site"
                                            191 DWFWCQAANN----ANVQHSALTVVPPGGD 216
                                                                        220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "cytoplasmic domain"
                                                              ..19
label= Signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                   'label= Mature protein
                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                            AAY95966 standard; Protein; 343 AA
                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    140..198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49..105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note=
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                                                                                                                                                                                                   (first
                                                                                                                                                                                                                               Human TANGO 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified-site
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                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                   05-DEC-2000
                                                                                                                                                                       AAY95966;
                                                                                                                                                                                                                                                                                                                                            Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                     Protein
                                                                                                                                                                                                                                                                                                                                                                                                                  Domain
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                                                                                                                 24
                                                                                                                 RESULT
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                                                                                                                                                                                                                                                        , human; bone marrow, umbilical cord venous endothelial cell; diagnosis; treatment.
94 EAHESGP----YKCKAQVTSCSKYSRDFSFTIVDPV-----TSPVLNIMVIQTETD 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       --LKSSPDLLTYFCRASSTSGAH------VDSARLQMHWELWSKPVSELRANFTLQD 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               138 RGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPC----HRQPANFSFL---PSQTS 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention provides polypeptides (AAY27127-Y27133) produced by human adult brain tissue, human bone marrow or a human umbilical cord venous endothelial cell. Host cells transformed with vectors comprising the nucleic acids encoding the polypeptides are used for the recombinant expression of the polypeptides. The polypeptides can be used in treatment and basic studies, with wide applications in treatment depending on the activity to be aimed at. Sequences AAX89112-125 represent nucleic acids encoding the polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37 EVFPKGRWVLITCCAPQPPPFITYSLCGTKNIKVAKKVVKTH------EPASFNLNVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44 KVVMKGQNVSMPCSHKNKSLQITYSLFRR------KTHPGTQDGKGEPAIFNLSIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.2%; Score 101.5; DB 20; Length 343; 26.2%; Pred. No. 0.24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New adult human brain tissue-produced polypeptides useful for
                                                                                                                                                                                                                             Human bone marrow-derived polypeptide (clone OAF038-Pro).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                     220
                                          FWCQAANN----ANVQHSALTVVPPGGD 216
                                                                                                                                                                                                                                                                                                                                           1..19
/note= "signal peptide"
                                                                                                                                                                                                                                                                                                                                                                    20..343
/note= "mature protein"
                                                            : |: | | : | | | | : |: | | | | YRCEAKNRL PNYATYSH-PVTMPSTGGD
                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 59-60; 86pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ξ
                                                                                                                                         AAY27130 standard; Protein; 343 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97JP-0358811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shibayama S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-419088/35.
N-PSDB; AAX89118, AAX89119.
                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ONOY ) ONO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              343 AA;
                                                                                                                                                                                                                                                        Brain tissue;
recombinant;
                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                               WO9933873-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fukushima D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-DEC-1998;
                                                                                                                                                                                                 14-SEP-1999
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                                          193
                                                                     194
                                                                                                                                                                       AAY27130;
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              142
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                                                                                                                                                                                                                                                                                                                                         Peptide
                                                                                                                                                                                                                                                                                                                                                                       Protein
                                                                                                                             AAX27130
                                                                                                             RESULT
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raise antibodies (useful as diagnostics) and to screen for modulator compounds.

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note= "tyrosine kinase phosphorylation site"
306..313
                                                                                                                                                                                                                               note= "tyrosine kinase phosphorylation site"
                                                                            269..272
/note= "casein kinase phosphorylation site"
288..291
"casein kinase phosphorylation site"
                           "casein kinase phosphorylation site"
                                                            'note= "casein kinase phosphorylation site"
                                                                                                                            "casein kinase phosphorylation site"
                                                                                                                                                               "casein kinase phosphorylation
                                                                                                                                                                                                                                                                                                                                                                                                  /note= "cell attachment site"
                                                                                                                                                                                                                                                                                                               274..279
/note= "N-myrisotylated"
293..298
                                                                                                                                                                                                                                                            /note= "N-myrisotylated"
                                                                                                                                                                                                                                                                                             'note= "N-myrisotylated"
                                                                                                                                                                                                                                                                                                                                                                 note= "N-myrisotylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                 'note= "leucine zipper"
            172..175
/note= "c
                                                                                                                              /note= "c
300..303
                                                                                                                                                               /note= "c
186..192
                                             217..220
                                                                                                                                                                                                                                                                                                                                                                                  266..268
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/note=
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                                         Modified-site
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WO200050443-A2

31-AUG-2000

25-FEB-2000; 2000WO-US05035

99US-0259387. 26-FEB-1999; (MILL-) MILLENNIUM PHARM INC.

Fraser CC;

WPI; 2000-533178/48. N-PSDB; AAA50441 Nucleic acids encoding TANGO 228, 240 and 243 pp. which have homology to the rat mast cell Ag-32, the Mycobacterium tuberculosis hypothetical protein Rv0712 and human phospholipase A2-activating protein

Claim 8; Fig 2; 188pp; English.

The present sequence is that of human TANGO 228, a protein that includes 2 Ig domains and which has homology to rat MCA-32 (mast call Ag-32), a cell surface antigen that is up-regulated in activated mast cells. The sequence was deduced from that of a cDNA clone (see AAA50441) isolated from a foetal spleen cDNA library. TANGO 228 proteins, nucleic acids and their modulators can be used to modulate the proliferation, differentiation and/or function of cells that form the spleen, e.g. to treat (foetal) spleen-associated diseases such as splenic lymphoma and/or splenomegaly, and/or phagocytic disorders such as those inhibiting macrophage engulfment of bacteria and viruses in the bloodsitram; to modulate mast cell function and thus to treat immunological disorders and diseases including allargic asthma and atopic dermatitis; to protect the body for treatment of anaphylactic shock or allergic dermatitis; to matchise type I immunological disorders, e.g. anaphylaxis or chinitis, by modulating the interaction between antigens and mast call receptors; and to treat tumour necrosis factor-related disorders (e.g. acute myocarditis, myocardial infarction, congestive heart failure); T cell disorders, and disorders related to anglogenesis (e.g. tumor formation, metastasis, cancer). TANGO 228 purpare, modulating the objection graceful and spottotic mediances, and disorders related complements of the content of expressed using gene therapy protocols. They can also be used to

2000US-225267P

14-AUG-2000;

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89 LKSSPDLLTYFCRASSTSGAH-----VDSARLOMHWELWSKPVSELRANFTLQDRG 139
                                                                                                                                                             36 HESGP----YKCKAQVTSCSKYSRDFSFTIVDPV-----TSPVLNIMVIQTETDR- 141
                                                                                                                                                                                           140 AGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPC----HRQPANFSFL---PSQTSDW 192
                                                                                                                                                                                                                ---HITLHCLSVNGSLPINYTFF----ENHVAISPAISKYDREPAEFNLTKKNPGEEEE- 193
                                                                                                    88
                                                                                                                          95
                                                                                                                        44 KVVMKGQNVSMFCSHKNKSLQITYSLFRRKTHLGTQDGK-----GEPAIFNLSITEA
                                                                                                   37 EVFPKGRWVLITCCAPQPPPITYSL-----CGTKNIKVAKKVVKTHEPASFNLNVT--
                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                      muscular disorder; reproductive disorder; gastrointestinal disorder; pulmonary disorder; cardiovascular disorder; renal disorder; proliferative disorder; inflammation.
                                                                              59;
                                                       Length 343;
                                                                                                                                                                                                                                                                                                                                                                                                           Human; gene therapy; neural disorder; immune system disorder;
                                                                             67; Indels
                                                     7.2%; Score 101.5; DB 21;
26.4%; Pred. No. 0.24;
Ive 27; Mismatches 67;
                                                                                                                                                                                                                                        216
                                                                                                                                                                                                                                                             194 YRCEAKNRLPNYATYSH-FVTMPSTGGD 220
                                                                                                                                                                                                                                        FWCQAANN----ANVOHSALTVVPPGGD
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                                                                                                                                                                                                                                                                                                                    ABB10224 standard; Protein; 362
                                                                 26.48;
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2000US-209467P.
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2000US-218290P.
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                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                             55; Conservative
                                                                                                                                                                                                                                                                                                                                                                                      Human cDNA SEQ ID NO: 532.
                                                                  Similarity
                                 343 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200154474-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-FEB-2000; 2
02-MAR-2000; 2
16-MAR-2000; 2
17-MAR-2000; 2
18-ARY-2000; 2
19-MAY-2000; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                               10-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JUL-2000;
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                                 Seguence
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                                                      Query Match
Best Local 8
                                                                             Matches
                                                                                                                                                                                                                                                                                               RESULT 25
88
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14-AUG-2000; 2000US-22526BP.

14-AUG-2000; 2000US-225270P.

14-AUG-2000; 2000US-225747P.

14-AUG-2000; 2000US-225757P.

14-AUG-2000; 2000US-225758P.

14-AUG-2000; 2000US-225758P.

14-AUG-2000; 2000US-225758P.

18-AUG-2000; 2000US-22678P.

22-AUG-2000; 2000US-22678P.

22-AUG-2000; 2000US-2268BP.

22-AUG-2000; 2000US-22894P.

30-AUG-2000; 2000US-229348P.

01-SEP-2000; 2000US-229348P.

01-SEP-2000; 2000US-229348P.

01-SEP-2000; 2000US-229348P.

01-SEP-2000; 2000US-229348P.

01-SEP-2000; 2000US-229348P.

06-SEP-2000; 2000US-229348P.

06-SEP-2000; 2000US-231414P.

08-SEP-2000; 2000US-231414P.

08-SEP-2000; 2000US-231414P.

08-SEP-2000; 2000US-231289P.

14-SEP-2000; 2000US-23139P.

14-SEP-2000; 2000US-23139P.

14-SEP-2000; 2000US-23139P.

14-SEP-2000; 2000US-23239P.

14-SEP-2000; 2000US-233306P.

25-SEP-2000; 2000US-233306P.

25-SEP-2000; 2000US-233306P.

27-SEP-2000; 2000US-233306P.

27-SEP-2000; 2000US-233306P.

28-SEP-2000; 2000US-233306P.

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20-SEP-2000; 2000US-233306P.

21-SEP-2000; 2000US-233306P.

21-SEP-2000; 2000US-233306P.

22-SEP-2000; 2000US-233306P.

23-SEP-2000; 2000US-233306P.

23-SEP-2000; 2000US-233306P.

23-SEP-2000; 2000US-233306P.

23-SEP-2000; 2000US-233306P.

23-SEP-2000; 200US-233306P.

23-SEP-2000; 2000US-233306P.

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29-SEP-2000; 2000US-236370P.
02-OCT-2000; 2000US-237033P.
02-OCT-2000; 2000US-237033P.
02-OCT-2000; 2000US-237033P.
02-OCT-2000; 2000US-237033P.
03-OCT-2000; 2000US-23993F.
03-OCT-2000; 2000US-249393P.
03-OCT-2000; 2000US-24188P.
03-OCT-2000; 2000US-24188P.
03-OCT-2000; 2000US-24180P.
03-OCT-2000; 2000US-24677P.
03-NOV-2000; 2000US-246477P.
03-NOV-2000; 2000US-246478P.
03-NOV-2000; 2000US-246478P.
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03-NOV-2000; 2000US-246478P.
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2000US-246526P
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The present invention provides human cDNAs, proteins and related genomic DNAs. These can be used in the treatment of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative disorders and inflammation. The present sequence is a protein of the invention.
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-----GEPAIFNLSITEA
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N-PSDB; ABA06446.
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                                             08-NOV-2000; 208-NOV-2000; 208-NOV-2000; 208-NOV-2000; 208-NOV-2000; 217-NOV-2000; 217
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115 HESGP----YKCKAQVTSCSKYSRDFSFTIVDPV-----TSPVLNIMVIQTETDR- 160

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05-SEP-2000; 2000US-0229509.
06-SEP-2000; 2000US-0229513.
06-SEP-2000; 2000US-0229513.
06-SEP-2000; 2000US-023913-1.
08-SEP-2000; 2000US-0231244.
08-SEP-2000; 2000US-0231244.
08-SEP-2000; 2000US-0231244.
08-SEP-2000; 2000US-0231414.
08-SEP-2000; 2000US-0231414.
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08-SEP-2000; 2000US-023399.
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13-OCT-2000; 2000US-023399.
13-OCT-2000; 2000US-023399.
13-OCT-2000; 2000US-023399.
13-OCT-2000; 2000US-02346178.
08-NOV-2000; 2000US-0246525.
08-NOV-2000; 2000US-0246528.
08-NOV-2000; 2000US-0246529.
08-NOV-2000; 2000US-02
     AGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPC----HRQPANFSFL---PSQTSDW 192
                                                                                                                                                                                                                                                                                                                                                Immunoglobulin; signal transduction pathway protein; cancer; antisense therapy; gene therapy; neurological disorder; renal disorder; carditovascular disorder; gastrointestinal disorder; pulmonary disorder; reproductive disorder; immune system disorder; proliferative disorder; muscular disorder.
                                                                                                                                                                                                                                                                                                                     Human immunoglobulin polypeptide SEQ ID No 163
                                                                                                * | : | | | | : | : | XRCEAKNRLPNYATYSH-PVTMPSTGGD 239
                                                                        FWCQAANN----ANVQHSALTVVPPGGD 216
                                                                                                                                                                                                      AAU18018 standard; Protein; 362
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2000US-0186564.
2000US-0189874.
2000US-0189874.
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2000US-020515.
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2000US-021886.
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used to diagnose a pathological condition or a susceptibility to a pathological condition or a subject by determining the presence or amount of amount of amount of amount of expression of the protein. Alternatively the identification of a binding partner to a sequence allows determination of changes in protein activity. The sequence callows determination of changes in receptors or other signal transduction pathway proteins that interact with the polypeptides of the invention and can be used to treat, prevent or diagnose various types of disorders such as neurological disorders, cardiovascular disorders, gastrointestinal disorders, muscular disorders, pulmonary disorders, pastrointestinal disorders, muscular disorders, pulmonary disorders, proliferative disorders and cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fite wipo.int/pub/published_pot_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated novel immunoglobulin polypeptide for monitoring the presence and progression of diseases and for diagnosis
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ive 27; Mismatches
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17-NOV-2000; 2000US-0249216.
17-NOV-2000; 2000US-0249217.
17-NOV-2000; 2000US-0249241.
17-NOV-2000; 2000US-0249245.
17-NOV-2000; 2000US-0249245.
17-NOV-2000; 2000US-0249265.
17-NOV-2000; 2000US-0249267.
17-NOV-2000; 2000US-0249297.
17-NOV-2000; 2000US-0251988.
0S-DEC-2000; 2000US-02519869.
0S-DEC-2000; 2000US-0251989.
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Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; imunosoppressive; antihiflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; crebroprotective; antiinflammatory; antiallacy; antidiabetic; antilloer; anticonvulsant; antifungal; antiparastic; cardiant; immune disorder; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine.
213 YRCEAKNRLPŅYATYSH-PVTMPSTGGD 239
                                                            ABP66811 standard; Protein; 362 AA
                                                                                                                       Human polypeptide SEQ ID NO 532.
                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-224519P.
2000US-225267P.
2000US-225270P.
2000US-22547P.
2000US-225757B.
2000US-225757B.
2000US-225757B.
2000US-229344P.
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                                                                                                    (first entry)
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                                                                                                    09-DEC-2002
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                                                                               ABP66811;
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Human protein sequence SEQ ID NO:1218.

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Novel polypeptide useful for diagnosis, prognosis, prevention, and
                                                                                                                                                        treatment of immune, hyperproliferative, renal, respiratory, cardiovascular, reproductive, endocrine, gastrointestinal and neurological disorders
                                                                                                                                                                                     Claim 11; SEQ ID NO 532; 369pp + Sequence Listing; English.
                                                                                                     Barash SC;
       20-OCT-2000; 2000US-241809P.
01-NOV-2000; 2000US-244617P.
17-NOV-2000; 2000US-249299P.
08-DEC-2000; 2000US-251868P.
08-DEC-2000; 2000US-251868P.
2000US-241785P
                                                                                                    Ruben SM,
                                                                                                                   WPI; 2002-681727/73.
N-PSDB; ABV83783.
                                                                                  (BARA/) BARASH S C.
                                                                (ROSE/) ROSEN C A. (RUBE/) RUBEN S M.
                                                                                                    Rosen CA,
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(ABP66710-ABP67129) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant) agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune through are menlytic ansemia, autoimmune thyroiditis, diabress mellitus, Crohn's disease, multiple sclerosis, rhebmatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing, (e) neurological diseases e.g. cerebral anoxia and parasitic infections. invention relates to novel genes (ABV83682-ABV84101) and proteins Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.

362 AA; Seguence

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13;
                                                                                                                                                                    LKSSPDLLTYFCRASSTSGAH------VDSARLQMHWELWSKPVSELRANFTLQDRG 139
                                                                                                                      63 KVVMKGQNVSMFCSHKNKSLQITYSLFRRKTHLGTQDGK-----GEPALFNLSITEA 114
                                                                                                                                                                                                                                         140 AGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPC----HRQPANFSFL---PSQTSDW 192
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                                                                                              37 EVFPKGRWVLITCCAPQPPPPITYSL-----CGTKNIKVAKKVVKTHEPASFNLNVT--
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                                                                59; Gaps
                    7.2%; Score 101.5; DB 23; Length 362; 26.4%; Pred. No. 0.26; tive 27; Mismatches 67; Indels 59;
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16-OCT-2001 (first entry)
            AAM25703;
HXXXH
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AAM25703 standard; Protein; 366 AA.

RESULT 26 AAM25703

13;

59; Gaps

67; Indels

27; Mismatches

55; Conservative

Matches

ð

88

37 EVFPKGRWVLITCCAPQPPPPITYSL-----CGTKNIKVAKKVVKTHEPASFNLNVT--

AAH99166 to AAH99904 encode the human proteins given in AAM25225 to AAM25963. The proteins can have activities based on the tissues and cells they are expressed in, such as: antiinflammatory; antihemmatic; ealls they are expressed in, such as: antiinflammatory; antihemmatic; cells they are expressive; antibacterial; endocrine; cardiant; central nervous system; virucide; antibacteric; vulnerary; cardiavascular; antianaemic; antialgery antiallergic; vulnerary; antidabetic; cytostatic; neuroprotective; antidepressant; nootropic; antidabetic; neuroprotective; antidabetic; sorcial gene therapy, antisense therapy and vaccine production; The proteins and polymucleotides are useful for screening for agonists or antagonists of a protein and for the treatment and diagnosis; of disorders associated with the activity of a protein e.g. inflammation, neuropathology, cardiac anaphylaxis, viral, bacterial, HIV and fungal infections, autoimmunity, genetic diseases, haematopoietic disorders, anaphylaxis, infections, wounds, burns, ulcers, and anaphylaxis, and anaphylaxis, whend the activity of a protein and fungal anaphylaxis, and anaphylaxis, whend and anaphylaxis, whend and anaphylaxis, whend and anaphylaxis, whend and anaphylaxis, thrombocytopaenia, wounds, burns, ulcers, and anaphylaxis, thrombocytopaenia, wounds, burns, ulcers, antiaggregant; haemostatic; vulnerary; antiulcer; osteopathic; eczema; dermatological; antiallergic; antiasthmatic; antidiabetic; cytostatic; neuroprocetive; antidepressant; noctropic; antiparkinsonian; infection; immunostimulant; gene therapy; antisense therapy; vaccine; inflammation; antianaphylactic; rheumatoid arthritis; septic shock; pancreatitis; cardiac dysfunction; neuropathology; cardiac anaphylaxis; autoimmunity; genetic disease; haematopoietic disorder; platelet disorder; asthma; thrombocytopaenia; osteoporosis; severe combined immunodeficiency; allergic rhinitis; diabetes; multiple sclerosis; depression; Alzheimer's disease; Parkinson's disease; neurodegenerative disorder; antiinflammatory, antirheumatic, antiarthritic; immunosuppressive; antibacterial; endocrine; cardiant; central nervous system; virucide; anti-HIV; fungicide; antimutagen; cardiovascular; antianaemic; anaemia; anaemia, platelet disorders, thrombocytopaenia, wounds, burns, ulcers, osteoporosis, severe combined immunodeficiency, eczema, allergic rhinitis, asthma, diabetes, cancer, multiple sclerosis, depression, Alzheimer's disease, Parkinson's disease, neurodegenerative and Isolated human polynucleotides encoding polypeptides, useful for the treatment and diagnosis of e.g. cancer, ulcers and HIV infection cancer; ulcer; HIV infection; human immunodeficiency virus; 7.2%; Score 101.5; DB 22; Length 366; 6.4%; Pred. No. 0.26; Claim 20; Page 252; 1217pp; English. rang YT, Liu C, Drmanac RT; 22-DEC-2000; 2000WO-US35017 23-DEC-1999; 99US-0471275. 21-JAN-2000; 2000US-0488725. 25-APR-2000; 2000US-0552317. 26.4%; neurological disorders. neurological disorder 2001-457603/49. (HYSE-) HYSEQ INC. Query Match Best Local Similarity 366 AA; N-PSDB; AAH99644 Homo sapiens. 26-JUL-2001. Sequence

RESULT 29

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ABG96265

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Homo sapiens.
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                                                                                                                                                                                 140 AGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPC----HRQPANFSFL---PSQTSDW 192
      --GEPAIFNLSITEA 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunoglobulin superfamily protein; IGSFP; cell proliferative disorder; atteriosclerosis; atherosclerosis; cirrhosis; hepatitis; paroxysmal nocturnal haemoglobinuria; polycythaemia vera; psoriasis; primary thrombocytopenia; cancer; leukaemia; lymphoma; sarcoma; myeloma; autoimmune disorder; inflammatory disorder; andemia; asthma; autoimmune dimnunodeficiency syndrome; AIDS; allergy; anaemia; asthma; autoimmune thyroiditis; contact dermatitis; crohn's disease; diabetes mellitus; glomerulonephritis; Godgasture's syndrome; gout; fraves' disease; Hashimoto's thyroiditis; irritable bowel syndrome; multiple sclerosis; osteoarthritis; osteoporosis; pancreatitis; Reiter's syndrome; theumatoid arthritis; Sjogren's syndrome; uveitis; trauma; viral infection; bacterial infection; fungal infection; herman, harman, 
                                                                                                                                                                                                                        ----VDSARLOMHWELWSKPVSELRANFTLQDRG
                                                                                                                       119 HESGP----YKCKAQVTSCSKYSRDFSFTIVDPV-----TSPVLNIMVIQTETDR-
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Forsythe IJ,
KVVMKGQNVSMFCSHKNKSLQITYSLFRRKTHLGTQDGK--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunoglobulin superfamily protein IGSFP-3.
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nn MR, Honchell CD, Burford N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG96265 standard; Protein; 385 AA.
                                                             LKSSPDLLTYFCRASSTSGAH ---
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21-SEP-2001; 2001US-323977P.
26-OCT-2001; 2001US-348447P.
02-NOV-2001; 2001US-343880P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-MAR-2002; 2002WO-US09052
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-MAR-2001;
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The present invention relates to new immunoglobulin superfamily proteins (IGSFP). The polypeptides, polynucleotides, agonists and antagonists are useful for diagnosing, treating or preventing disorders associated with

Yue H,

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ateriosclerosis, atherosclerosis, cirrhosis, hepatitis, paroxysmal nocturnal haemoglobinuria, polycythaemia vera, psoriasis, primary thrombocytopenia or cancers including leukaemia, lymphoma, sarcoma or myeloma, and autoimmune or inflammatory disorders, e.g. acquired immunodeficiency syndrome (AIDS), allergies, anaemia, asthma, autoimmune thyroiditis, contact dermaitis, Crobm's disease, diabetes mellitus, glomerulonephritis, Goodpasture's syndrome, gout, Graves' disease, lashinoc's thyroiditis, irritable bowel syndrome, multiple sclerosis, osteoarthritis, osteoporosis, pancreatitis, Reiter's syndrome, creatial, fungal, parasitic, protozoal or helminthic infections. They are also useful in the assessment of the effects of exogenous compounds on the expression of nucleic acid and amino acid sequences of ISSPP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     138 HESGP----TSPVLNIMVIQTETDR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              37 EVFPKGRWVLITCCAPQPPPFITYSL-----CGTKNIKVAKKVVKTHEPASFNLNVT--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LKSSPDLLTYFCRASSTSGAH-----VDSARLQMHWELWSKPVSELRANFTLQDRG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             59; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                     DB 23; Length 385;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CD31; domain; antibody; detection; carcinoma; inflammation; inhibition; treatment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Methods of screening for inhibitors of CD31 interactions mapping their sites of reaction with the CD31 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 7.2%; Score 101.5; DB 23; Best Local Similarity 26.4%; Pred. No. 0.28; Matches 55; Conservative 27; Mismatches 67;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR94894 standard; peptide; 571
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CD31 fragment (domains D1-D6)
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                                                                                                                                                                                                                                                                                                                                                                         385 AA;
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13

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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                               438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         326 NFTIQKEDTIVSQTQDF---TKIASKSDSGTYICTAGIDKVVKKSNTVQIVVCEMLSQPR 382
                                                                                                                                                                                                                                                                                                                                                                                           127 SELRANFTLODRGAGPRVEMICOASSGSPPITNSLIGKDGOVHLOORPCHROPANFSFLP 186
                                                                                                                                                                                                                                                                                             275 KVESSRISKVSSIVVNITELFSKPELESSFTHLDQGERINLSCSIFGAPP-----A
                                                                                                                                                                                                                                                                           ----GRWVLITCCAPQPPPPITYSLCGTK
                                                                                                                                                                                                                                                                                                                                                                                                            383 ISYDAQFEVI---KGQTIEVRCESISGTLPISYQLL-KTSKVLENSTKNSNDPAVFKLNP
                        Screening of inhibitors of CD31 is achieved by incubating labelled CD31 component with potential inhibitor, adding this mixture to CD31 component immobilised on a support, washing and detecting label. Alternatively, potential inhibitor can be incubated with CD31 component immobilised on a support and labelled CD31 component can added followed by washing and detecting label. Failure to detect label suggests that the compound being screened is not an inhibitor of CD31. The method is used to identify antibodies that can be used in the treatment of carcinomas and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin superfamily receptor translocation associated; IRTA;
IRTA2b; human; immunoglobulin receptor; Fc receptor; melanoma;
lymphoma; myeloma; B cell malignancy; cancer; chromosome 1q21;
                                                                                                                                                                                                                                                68;
                                                                                                                                                                                                            7.2%; Score 101; DB 17; Length 571; 21.2%; Pred. No. 0.55; -ive 39; Mismatches 116; Indels 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GSGPITYKFYREKEGKPFYQMTSNATQAFWTKQKASKEQEGEY 540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WQGPLESPIL----ALPLYRST-----RRLSEEFGGF 250
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/note= "Asn is N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    383..385
/note= "Asn is N-glycosylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunoglobulin receptor isoform IRTA2b.
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/label= Signal_peptide
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/label= Mature_protein
Disclosure; Figure 16; 31pp; English.
                                                                                                                                                                                                                                                                      19 KAREESITPVVSIAYKVLEVFPK---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB82314 standard; Protein; 592 AA
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                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis; therapy
                                                                                                                                                                                571 AA;
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                                                                                                                                                                                                                                           90;
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                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           125 PVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPAN--- 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ---FSF-LPSQTSDWFWCQAANNANVQHS----ALTVVPPGGDQKMEDWQGPLESPILA 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         380 PVLNLSSPEDLIFEGA--KVTLHCEÁQRGSLPILY-----ÓFHHEDAALERRSANSAG 430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ---PVSHPVLT 476
                                                                                                                                                                                                                                                                                                                                                    The present sequence is that of the mover numentaglobulin.

Treceptor, immunoglobulin apperfamily receptor translocation
associated protein isoform 20 (IRTA2b), an Fc receptor involved in
the pathogenesis of lymphoma and melanoma. Effecting band 1921 in
the pathogenesis of lymphoma and melanoma. Effecting band 1921 in
multiple myeloma and B cell lymphoma led to the discovery of IRTA2

and IRTA1 (see ABB2212) as founding members of a novel subfamily
of related receptors within the immunoceoptor family. The IRTA2

clocus is transcribed into 3 major mRNA isoforms, IRTA2a in IRTA2

clocus is transcribed into 3 major mRNA isoforms, IRTA2a in IRTA2

clocus is transcribed into 3 wajor mRNA isoforms, IRTA2a in IRTA2

clocus is transcribed into 3 vesiques, whose hydrophobicity suggest
did plycoprotein, which diverges from IRTA2a at residue 560,

extending for a further 32 residues, whose hydrophobicity suggest
display a specific pattern of expression in mature B cells. The IRTA2

is expressed in GC centrocytes and in perifollicular cells, which may
include immunoblasts and memory cells. The invention provides IRTA

include addis and proteins, and antibodies directed to an epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             malignancy comprising a lg21 chromosomal rearrangement using a malignancy comprising a lg21 chromosomal rearrangement using a nucleic acid molecule that specifically hybridises with a unique sequence of human IRTA-5, and treating a subject having a B cell cancer by administering an anti-IRTA antibody or an antisense oilgonucleotide that specifically hybridises to IRTA mRNA so as to prevent overspression of IRTA protein and hence to arrest coll growth or induce cell dath of cancer cells expressing IRTA.

The B cell cancer is selected from B cell lymphoma, mantle cell lymphoma, miltiple myeloma, Burkitt's lymphoma, marginal zone lymphoma, diffuse large cell lymphoma and follicular lymphoma. The B cell lymphoma is selected from mucosa-associated-lymphoma. The B cell lymphoma or non-Hodgkin's lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               67 --NIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                            New genes encoding immunoglobulin receptor, Immunoglobulin super
Receptor Translocation Associated proteins, used to treat B cell
                                                                                                                                                                                                                                                                                                                                            present sequence is that of the novel human immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.2%; Score 101; DB 22; Length 592; 22.8%; Pred. No. 0.58; ve 33; Mismatches 89; Indels 6
                                                                                                                                                                                                                                               malignancies including lymphomas and multiple myeloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       431 GVAISFSLTAEHSGNYYCTADNGFGPQRSKAVSLSITV----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 SIAYKULEVFPKGRWVLITCCAPQP----PPPITYSLCGTK
                                                                                                                                                                                                                                                                                            Claim 3; Fig 18B-1-18B-2; 72pp; English.
                           (UYCO ) UNIV COLUMBIA NEW YORK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22.8%;
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                                                                                                                           WPI; 2001-355921/37
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Best Local Similarity
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                                                                                                                                                  N-PSDB; AAF30951
                                                                         Dalla-Favera R;
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Secreted protein, SECP; human, antiarteriosclerotic; antiatherosclerotic; antiatherosclerotic; antiatherosclerotic; antiatherosclerotic; antiantiabelic; antiantial manatory; neuroprotective; antialcer; antiantiamento; antialcer; antiantemento; antialcer; antiantemento; antialcer; antiantic; hypotensive; gonadal dysgenesis; antiparkinsonian; ophthalmological; immunosuppressive; perioarditis; antiparkinsonian; ophthalmological; carrhosis; hepatitis; angina pectoris; autoimmune disorder; AIDS; Crohn's disease; multiple sclerosis; epilepsy; lucerative collits; cardiovascular disorder; myocardial infarction; Huntington's disease; Alzheimer's disease; Creutzfeldt-Jakob disease; developmental disorder; buchenne muscular dystrophy; antipsoriatic;
                                                                                Human SECP-20 protein from clone 8039739CD1 SEQ ID 20.
                                                                                                                                                                                                                                                          Becker muscular dystrophy; Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 75; Page 165-166; 188pp; English.
            ABB84668 standard; Protein; 727 AA.
                                                                                                                                                                                                                                                                                                                                                                                             06-APR-2001; 2001US-282112F.
09-APR-2001; 2001US-282702F.
13-APR-2001; 2001US-283855F.
19-OCT-2001; 2001US-343718F.
07-DEC-2001; 2001US-339256F.
13-FEB-2002; 2002US-357002F.
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Gietzen KJ, Gri
ee EA, Lee S, Lu
t, Ramkumar J, T
Yao MG, Xu Y,
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                                                            (first entry)
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Swarnakar A, Kamoo...
Swarnakar A, Yao MG,
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                                                                                                                                                                                                                                                                                                         WO200279441-A2.
                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                30-MAR-2001;
                                                           11-FEB-2003
                                                                                                                                                                                                                                                                                                                                  10-0CT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gandhi AR,
                                   ABB84668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lal PG,
ABB84668
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ason PM, Sanjanwala MM; Thangavelu K, Tran UK, Walia NK; Forsythe IJ; Ison CH; Elliott VS, Honchell CD, Duggan BM, E JA, He A, E , Mason PM, Lu DAM, Ma Tang YT, ', Yue H; Ding L, Dug Griffin JA,

Novel human secreted protein useful for treating, preventing or diagnosing cancer, hepatitis, psoriasis, asthma, diabetes mellitus, anemia, epilepsy, cataract, Alzheimer's disease

antiarteriosclerotic, antiatherosclerotic, hepatotropic, cytostatic, antialteriosclerotic, antiatherosclerotic, antianaemic, antidiabetic, antiallematory, neuroprotective, antibloer, antidiabetic, vasotropic, antiarthritic, cardiant, hypotensive, anticonvulsant, nootropic, immunosuppressive, antiparkinsonian and ophthalmological activity. The polynucleotides and polypeptides of the invention can be used for diagnosing, treating or preventing cell proliferative disorder e.g. arteriosclerosis, atherosclerosis, cirrhosis, hepaticis, cancer, autoimmune/inflammatory disorders e.g. acquired immunodeficiency syndrome (AIDS), allergies, asthma, anaemia, diabetes mallitus, crohn's disease, multiple sclerosis, ulcerative colitis, posriaais, theumatoid arthritis, etc. cardiovascular disorder e.g. myocardial infarction, angina pectoris, hypertension, Raymaud's disease, myocarditis, pericarditis, etc. churchogical disorders e.g. epilepsy, Huntington's disease, Parkinson's disease, Alzheimer's disease, Creutzfeldt-Jakob disease, petc; and developmental disorders e.g. Duchenne and Becker muscular dystrophy, This invention describes novel secreted proteins (SECP) which have

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11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cataract, gonadal dysgenesis, Cushing's syndrome, etc. The products of the invention can also be used for drug screening, proteome analysis, microarrays creating knock-in humanised animals or transgenic animals to model human diseases, in somatic or germline gene therapy, to generate a transcript image of a tissue or cell type, for detecting differences in the chromosomal location due to translocation, inversion, etc., among normal, carrier or affected individuals, and as hybridization probes for mapping naturally occurring genomic sequences. ABB84649-ABB84673 represent secreted proteins encoded by the cDNA's shown in ABS57545-ABS57569, described in the disclosure of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                      367 VIVRIPVSHPVLTFRAPRAHTVVGDLLELHCESLRGSPPILYRFY-HEDVTLGNSSAPSG 425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    79 EPASFNLNVTLKSSPDLLTYFCRASSTSGA-HVDSARLOMHWELWSKPVSELRANFTLQD 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RGA----GPRVEMICQASSGSPPITNSLIGKD---GQVHLQQRPCHRQPANFSF-LPSQT 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  476 PGAQAVVGDLLELHCESLRGSFPILYWFYHEDDTILGNISAHS----GGGASFNISLTTEH 531
                                                                                                                                                                                                                                                                                                                                                                                           25 ITPVVSIAYKVLEV-FPKGRWVL----ITCCAPQPPPJITYSLCGTKNIKVAKKVVKTH 78
                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                        24; Length 727;
                                                                                                                                                                                                                                                                                                   7.1%; Score 100; DB 24; Length 7
4.9%; Pred. No. 0.97;
ve 39; Mismatches 88; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SDWFWCQAANNANVQHSALTVVPPGGDQK 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  532 SGNYSCEADNGLGAQHSKVVTLNVTGTSR 560
                                                                                                                                                                                                                                                                                                                             24.98;
                                                                                                                                                                                                                                                                                                                        Local Similarity 24.9%
nes 52; Conservative
                                                                                                                                                                                                                                                     727 AA;
                                                                                                                                                                                                                                                          Sequence
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RESULT 33 AAB82316

AAB82316 standard; Protein; 734 AA.

AAB82316;

(first entry) 23-JUL-2001

Human immunoglobulin receptor IRTA3 protein.

Immunoglobulin superfamily receptor translocation associated;
IRTA3; human; immunoglobulin receptor; Fc receptor; melanoma;
iymphoma; myeloma; B cell malignancy; cancer; chromosome 1921;

Homo

diagnosis; therapy

WO200138490-A2.

31-MAY-2001

28-NOV-2000; 2000WO-US32403.

29-NOV-1999;

(UYCO) UNIV COLUMBIA NEW YORK,

Dalla-Favera R;

WPI; 2001-355921/37. N-PSDB; AAF30953.

New genes encoding immunoglobulin receptor, Immunoglobulin super Receptor Translocation Associated proteins, used to treat B cell malignancies including lymphomas and multiple myeloma -

Claim 4; Fig 18C-1-18C-2; 72pp; English

The present sequence is that of the novel human immunoglobulin

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pathogenesis of lymphoma and melanoma. Efforts to identify genes involved in the pathogenesis of lymphoma and melanoma. Efforts to identify genes in chromosomal aberrations affecting band lg21 in multiple mytholved in chromosomal aberrations affecting band lg21 in multiple mytholved in chromosomal aberrations affecting band lg21 in multiple mytholwed in chromosomal led to the discovery of IRTA1 and IRTA2 (see AAB82312-15) as founding members of a novel subfamily of related receptors within the immunoreceptor family. 3 Additional proteins, IRTA3 in RTA4 and IRTA5 (see AAB82316-18), were subsequently the immunoreceptor family. The IRTA3 is expressed in GC centrocytes and in perifolicular cells. Which may include lymphoblasts and memory cells. This is call analogous to IRTA2 expression. The invention provides IRTA nucleic acids and proteins, and antibodies directed to an epitope of an IRTA protein. Methods are claimed for: detecting a B cell malignancy comprising a 1q21 chromosomal rearrangement using a nucleic acid molecule that specifically hybridises with a unique sequence of human IRTA1-5, and treating a subject having a B cell malignancy concernitionally hybridises to IRTA mand so as to prevent concerned from B cell sexpressing IRTA. The B cell additional participal and follicular imphoma, mantle cell lymphoma, diffuse large cell lymphoma, mantle cell lymphoma, the B cell lymphoma is selected from mucosa-associated-lymphoma. The B cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RGA----GPRVEMICQASSGSPPITNSLIGKD---GQVHLQQRPCHRQPANFSF-LPSQT 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      483 PGAQAVVGDLLELHCESLRGSFPILYWFYHEDDTLGNISAHS----GGGASFNLSLTTEH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        79 EPASFNINVTLKSSPDILITYFCRASSTSGA-HVDSARLQMHWELWSKPVSELRANFTLQD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25 ITPVVSIAYKVLEV-FPKGRWVL----ITCCAPQPPPPITYSLCGTKNIKVAKKVVKTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antianginal; gene therapy;
heart failure; angina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.1%; Score 100; DB 22; Length 734; 24.9%; Pred. No. 0.99;
   immunoglobulin superfamily receptor translocation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              39; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      190 SDWFWCQAANNANVQHSALTVVPPGGDOK 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SGNYSCEADNGLGAQHSKVVTLNVTGTSR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABG74786 standard; Protein; 31267 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RGS11; human; screening; cardiant;
heart disorder; cardiac ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2002; 2002WO-JP06019.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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es 52; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     734 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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Matches
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6398 ELKPVEVVKYSDVE-----LECEVTGTPPFEVTWLKNNREIRSSKKYTLTDRVSVF 6448
                                                                                                                                                                                                                                   This invention describes a novel method for screening compounds for their ability to regulate the activity and expression of human RGSI1 and its partial peptides and salts, by observing the expression or activity of RGSI1 in the presence or absence of the test compound. The products of the invention have cardiant and antianginal activity and can be used for the invention have cardiant and antianginal activity and can be used for given thearpy. The methods and compositions are useful in the prevention, treatment and diagnosis of heart disorders such as cardiac ischaemia, heart failure and angina. This sequence represents the human RGSI1 protein described in the disclosure of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6449 NLHITKCDPSDIGEYQCIVSNEGGSCSCSTRVAL----KEPPSFIKKIENTITVLKSSA 6503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      multiple sclerosis, diabetes; genetic disorder; wound; burn; infection; arthritis; cytostatic; immunomodulator; nootropic; neuroprotective; antiparkinsonian; antidiabetic; immunosuppressive; dermatological; haemostatic; vulnerary; fungicide; antibacterial; virucide; protozoacide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24 EITPVVSIAYKVLEVFPKGRWVLITCCAPQPPPPITYSLCGTKNIKVAKKVVKTHEPASF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ----SEVDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TLQDRGAGPRVEMICQASSGSPPIT-----NSLIGKDGQVHLQQRPCHRQPANFSFLPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, genome mapping, gene therapy, food supplement; virus; fungus, cell-proliferative disorder; neurodegenerative disease; bacterial, Parkinson's disease; Alzheimer's disease; autoimmune disease;
                                                                                                                                                     expression and activity for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 99.5; DB 24; Length 31267;
Pred. No. 2.3e+02;
1; Mismatches 72; Indels 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ----GSPPISITWLKDDQILDEDDNVYI---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84 NLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSKPVS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6540 VATLQIRSVDNGHSGRYTCQAKNESGVERCYAFLLVOEP 6578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     --SDWFWCQAANNANVQ--HSALTVVPP 213
                                                                                                                                             Screening compounds regulating RGS11 expre
prevention and treatment of heart disease
                                                                                                                                                                                                    Claim 1; Page 59-261; 321pp; Japanese.
                                                      Yamamoto K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human polypeptide SEQ ID NO 1330.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABP69283 standard; Protein; 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.18; 22.48;
               (TAKE ) TAKEDA CHEM IND LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAR-2002; 2002WO-US05095.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        49; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                      Tanida S,
                                                                                                                                                                                                                                                                                                                                                                                                                              31267 AA;
                                                                                     WPI; 2003-167557/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                            N-PSDB; ABX13540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6504 TFOSTVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     188 QT----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200270539-A2
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                                                    Koyama N,
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP69283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 35
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Homo sapiens.
   The invention relates to an isolated polymucleotide (I) comprising a nucleotide sequence selected from any of 948 sequences

(ABZI119-ABZI2066) or their mature protein coding portion, active domain coding protein or complementary sequences. The polymucleotides are useful coding protein or complementary sequences. The polymucleotides are useful coding protein or complements sequences. The polymucleotides are useful considered polypeptides (ABBE68902-ABP69849) are useful as molecular weight markers, as food supplement, for generating antibodies, in medical imaging, screening and diagnostic assays and for treating coll-proliferative disorders (cancer), neurodegenerative diseases

(Parkinson's or Alzabeimer's disease), autoimmune diseases (multiple sclerosis, diabetes, lupus) genetic disorders, myeloid or lymphoid disorders, platelet or coagulation disorders, wound, burns, incision, parasitic), arthritis, etc.

Darasitic), arthritis, etc.

Darasitic), arthritis, etc.

Darasitic), arthritis, etc.

Specification, but was obtained in electronic format directly from WIPPO set fip.wipo.int/pub/published_pot_sequences.
                                                                                                                                                                                                                                            domain
useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         --NIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSK 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PVSELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPAN--- 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PVINLSSPEDLIFEGA--KVTLHCEAQRGSLPILY-----QFHHEDAALERRSANSAG 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    73
                                                 Ren F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 TMPYSVISDSPRS-WIQVQIPASHPVLTLSPEKALNFEGTKVTLHCETQEDSLRTLYRFY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       74 HEGVPLRHKSVRCERGASISFSLTTENSGN---YYCTADNGLGAKPSKAVSLSVTVPVSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                            New polynucleotides comprising sequences assembled from expressed sequence tags (ESTs), useful for treating cell-proliferative, neurodegenerative, autoimmune, genetic, myeloid or lymphoid, or platelet or coagulation disorders
                                               Zhao QA, 1
Ghosh M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 222;
                                                                                                                                                                                       Claim 9; SEQ ID NO 1330; 1012pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     86; Indels
                                            Goodrich RW, Asundi V, Zhang J, Z
Aa Y, Yamazaki V, Chen R, Wang Z,
, Wang D, Drmanac RT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30 SIAYKULEVFPKGRWVLITCCAPQP----PPPITYSLCGTK
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 7.0%; Score 98.5; DB 23; Best Local Similarity 23.9%; Pred. No. 0.25; Matches 50; Conservative 28; Mismatches 86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GVAISFSLTAEHSGNYYCTADNGFGPORS
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 05-MAR-2001; 2001US-0799451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (updated)
(first entry)
                                             Zhou P, Gc
Yang Y, Ma
T, Wang J,
                                                         Ma
                                                                                          WPI; 2002-759812/82.
                       (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                     222 AA;
                                                                                                       N-PSDB; ABZ11500
                                                         Xue AJ, Ya:
Wehrman T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
10-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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PECAM-1 and variants can be used in modulating angiogenic processes which depend on neutrophil chemotaxis and/or formation of junctions between endothelial calls, e.g. in tunnour development. PECAM-1, variants and antibodies to them can be used in the prevention of metastatic disease. The antibodies can also be used to produce anti-indiotypic antibodies which can be used to sequester anti-PECAM antibodies in an individual, thereby to treat or prevent pathological conditions which may be associated with an immune response whereby PECAM-1 is recognised as foreing by the immune system of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cysteine residues, spaced approx. 50 amino acids apart throughout the entire external domain of PECAM-1, are thought to participate in disulfide-bond formation within individual immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprises the N-terminal sequence QENSF and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Platelet and endothelial cell adhesion molecule - the molecule its antibodies, are useful for preventing metastatic disease
                                                                                                                                   602..620
/label= transmembrane_domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The variant pref. comprises the N-terminal ses 574 amino acids long. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= phosporylation_site
                                                                                                                                                                                                                  label= glycosylation_site
                                                                                                                                                                                                                                                                         'label= glycosylation_site
                                                                                                                                                                                                                                                                                                                                 /label = glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                     /label= glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                                                                             /label= glycosylation_site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= glycosylation_site
                                                                           28..738
/label= mat_protein
                                                 label= sig_peptide
location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Fig 1; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BLOO-) BLOOD CENT SE WISCO.
(NEWM/) NEWMAN P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90WO-US07418
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                                                                                                                                                                                     Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-DEC-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Newman PJ;
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                    Peptide
                                                                              Protein
                                                                                                                                   Domain
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Score 98; DB 12; Length 738; Pred. No. 1.6;

7.0%;

Best Local Similarity

Query Match

Platelet and endothelial cell adhesion molecule, antibodies, mutein; leukemia; cancer; variant.

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seck P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 38
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  11;
                                               67 NIKVAKKVVKTHEPASFNINVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWELWSKPV 126
                                                              356 NFTIQKEDTIVSQTQDF---TKIASKSDSGTYICTAGIDKVVKKSNTVQIVVCEMLSQPR 412
                                                                               127 SELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANFSFLP 186
                                                                                                                         TEDVE-YQCVADNCHSHARMLSEVLRVKVIAPVDEVQISILSSKVVESGEDIVLQCAVNE 527
                                  -----A 355
                                                                                         -- VQHSALT--- VVPPGGDQKME----D 221
                  99
               19 KARBEEITPVVSIAYKVLEVFPK------GRWVLITCCAPQPPPITYSLCGTK
   Gaps
  68;
                                                                                                                                                                                                                                                                     PECAM-1; platelet/endothelial cell adhesion molecule-1; CD31; inhibitor; allergy; adult respiratory distress syndrome; Crohn's disease; septic shock; traumatic shock;
                                                                                                                                                                                                                                                                                             multi-organ failure; autoimmune disease; asthma; inflammatory bowel disease; psoriasis; rheumatoid arthritis; reperfusion injury; stroke; cancer; multiple sclerosis; atherosclerosis; leukaemia; organ transplant; therapy.
                              305 KVESSRISKVSSIVVNITELFSKPELESSFTHLDOGERLNLSCSIPGAPP
                                                                                                                                                         528 GSGPITYKFYREKEGKPFYQMTSNATQAFWTKQKASKEQEGEY 570
                                                                                                                                              222 WQGPLESPIL----ALPLYRST------RRLSEEEFGGF 250
 116;
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                       'label= Extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                602..620
/label= Transmembrane
                                                                                                                                                                                                                                                                                                                                                                              28..738
/label= Mat_protein
                                                                                                                                                                                                                                                                                                                                                                                                                               621..738
/label= Cytoplasmic
                                                                                                                                                                                                                                                                                                                                                                       'label= Sig_peptide
                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                      standard; Protein; 738
39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label= Ig-like
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249..339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       label= Ig-like
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424..515
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/label= Ig-like
                                                                                                               SQTSDWFWCQAANNAN----
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                                                                                                                                                                                                                                       entry)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ..423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                145..248
                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9710839-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-SEP-1996;
                                                                                                                                                                                                                                      17-0CT-1997
90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAR-1997
                                                                                                               187
                                                                                                                              469
                                                                                                                                                                                                      AAW14802
                                                                                                                                                                                                                      AAW14802;
                                                                                                                                                                                                                                                      PECAM-1.
                                                                                                                                                                                                                                                                                                                                                      Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                              Protein
                                                                                                                                                                                                                                                                                                                                                                                               Domain
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Matches
                                                                                                                                                                                      RESULT 37
                                                                                                                                                                                               AAW14802
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356 NFTIOKEDTIVSOTODF---TKIASKSDSGTYICTAGIDKVVKKSNTVQIVVCEMLSQPR 412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ----VOHSALT--VVPPGGDOKME----D 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  469 TEDVE-YQCVADNCHSHAKMLSEVLRVKVIAPVDEVQISILSSKVVESGEDIVLQCAVNE 527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -GRWVLITCCAPOPPPITYSLCGTK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      67 NIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSKPV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PECAM-1 (AAW14802), or platelet/endothelial cell adhesion molecule-1 or CD31, is a glycoprotein that is constitutively expressed on the surface of endothelial cells, platelets and most leukocytes. It recognises and binds to other PECAM-1 molecules present on the surface of adjacent cells. Novel, synthetic linear and cyclic peptides (AAW28157) of 4-13 amino acids are based on the C2-type Ig domain of PECAM-1. They inhibit the binding of PECAM-1 to itself and can be used for treating allergy, adult respiratory distress syndrome. Crohn's disease, septic shock, milti-cronn failure surformment disease actions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; platelet-endothelial cell adhesion molecule-1; PECAM-1; angiogenesis; inflammation; arterial occlusion; tumour development;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 traumatic shock, multi-organ failure, autoimmune disease, asthma, finflammatory bowel disease, psoriasis, rheumatorid arthritis, reperfusion injury, stroke, cancer, organ transplants, multiple sclerosis, atherosclerosis and leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    New synthetic peptide(s) based on PECAM-1 sequences - used for inhibiting the binding of PECAM-1 to itself, for treating e.g. asthma, autoimmune diseases, shock, strokes or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 7.0%; Score 98; DB 18; Length 738; Best Local Similarity 21.2%; Pred. No. 1.6; Matches 60; Conservative 39; Mismatches 116; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ----RRLSEEFGGF 250
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                                                                                                                                                                                                                                                                                             Revelle BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A platelet-endothelial cell adhesion molecule-1.
                                                                                                                                                                                                                                                                                             Ren K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 31-34; 87pp; English.
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                                                                                                                                                                                                                           BIOTECHNOLOGY CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   222 WQGPLESPIL----ALPLYRST-
                                                                                                                                                                                                                                                                                         Kint PN,
                 950S-0003941.
950S-0003951.
950S-0003953.
950S-0003968.
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95US-0003996
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                                                                                                                                                                                                                                                                                      Bjercke RJ,
S:
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                                                                                                                                                                                                                        (TEXA-) TEXAS
                        19-SEP-1995;
19-SEP-1995;
19-SEP-1995;
                                                                                                                    19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-NOV-2000
                                                                                                                                                                                                                                                                                                                        Sherwood
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11;

Fri Feb

(first entry)

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standard; Protein; 738 AA.
                      AAB65866
                                          AAB65866;
RESULT 39
          AAB65866
ID AAB6
XX
AC AAB6
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Human; mouse; secreted protein; TANGO253; TANGO 257; TANGO 281;
TYTRRCEPT 758; coronary disorder; olfactory disorder;
neurological disorder; pulmonary disorder; immunological disorder;
developmental disorder; kidney disorder
                                                                                                                                                                                                                                                                                                                                                                             New nucleic acids for treating diseases and disorders, e.g. atherosclerosis, infection, autoimmune diseases, obesity, ear disorders, brain disorders, tumors, diabetes, arthritis, multiple sclerosis and asthma
                                                 Human PECAM-1 protein SEQ ID NO: 73.
                                                                                                                                                                                                                                                                                         (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                    19-JUN-2000; 2000WO-US16883.
                                                                                                                                                                                                                                                                                                                      Leiby KR, McKay C,
                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-050109/06.
                                                                                                                                                                            WC200078808-A1
                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                              18-JUN-1999;
                      28-MAR-2001
                                                                                                                                                                                                         28-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 67 NIKVAKKVVKTHEPASFNLAVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWELWSKPV 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             356 NFTIQKEDTIVSQTQDF---TKIASKSDSGTYICTAGIDKVVKKSNTVQIVVCEMLSQPR 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ---VQHSALT--VVPPGGDQKME----D 221
                                                                                                                                                                                                                                                                                                             Therapeutic methods useful for modulating angiogenic processes, relieving inflammation, or inhibiting arterial occlusions by administering a soluble form of the platelet-endothelial cell adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    305 KVESSRISKVSIVVNITELFSKPELESSFTHLDQGERINLSCSIPGAPP-----A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        127 SELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANFSFLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 KAREEEITPVVSIAYKVLEVFPK------GRWVLITCCAPQPPPPITYSLCGTK
                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a human platelet-endothelial cell adhesion molecule-! (PECAM-1) polypeptide. A soluble form of PECAM-1 is used for modulating angiogenic processes; relieving inflammation, and inhibiting arterial occlusions. The method is useful for modulating angiogenic processes that are associated with tumour development, relieving inflammation due to leukocyte transmigration (e.g. arthritis, bee sting, spider bite, sepsis or anaphylactic shock), or inhibiting arterial occlusions that are associated with atherosclerosis or vascular trauma. PECAM-1 isoforms are useful for making antibodies, e.g. monoclonal antibodies, for various diagnostic and therapeutic uses.
  spider bite; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 7.0%; Score 98; DB 21; Length 738; Best Local Similarity 21.2%; Pred. No. 1.6; Matches 60; Conservative 39; Mismatches 116; Indels (
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 leukocyte transmigration; arthritis; bee sting; spid
anaphylactic shock; atherosclerosis; vascular trauma
                                                                                                                                                                                                                CENT SOUTHEASTERN WISCONSIN.
                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Column 37-42; 22pp; English.
                                                                                                                                                                                                                                              PJ;
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                                                                                                                              95US-0478208,
                                                                                                                                                           90US-0466140.
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                                                                                                                                                                                                                                             Gumina RJ,
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                                                                                                                                                                                                                (BLOO-) BLOOD
                                           Homo sapiens
                                                                                                                                                                                                                                          Kirshbaum N,
                                                                                                                                                                                    16-NOV-1994;
                                                                                                  11-JUL-2000
                                                                      US6087331-A
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e.g. ear

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                                           The present invention provides the protein and coding sequences of the human and murine secreted or transmembrane proteins TANGO 253, TANGO 257, TANGO 281 and INTERCEPT 258. These are useful in the treatment of coronary, pulmonary, olfactory, immunological, neurological, developmental and kidney disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          469 TEDVE-YQCVADNCHSHAKMLSEVLRVKVIAPVDEVQISILSSKVVESGEDIVLQCAVNE 527
                                                                                                                                                                                                                                                                                                                                                    99
                                                                                                                                                                                                                                                                                                                                                    ---GRWVLITCCAPOPPPITYSLCGTK
                                                                                                                                                                                                                                                                                                                                                                                                                                                 67 NIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLQMHWELWSKPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                356 NFTIQKEDTIVSQTQDF---TKIASKSDSGTYICTAGIDKVVKKSNTVQIVVCEMLSQPR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              127 SELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANFSFLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               413 ISYDAQFEVI --- KGQTIEVRCESISGTLPISYQLL-KTSKVLENSTKNSNDPAVFKDNP
                                                                                                                                                                                                                                                                                                   68;
                                                                                                                                                                                                                                                                                                                                                                                 | | | | : :: | | | | 305 KVESSRISKVSSIVVNITELFSKPELESSFTHLDQGERLNLSCSIPGAPP-
                                                                                                                                                                                                                                                  DB 22; Length 738;
                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.0%; Score 98; DB 22; Le
21.2%; Pred. No. 1.6;
cive 39; Mismatches 116;
Disclosure; Page 255-258; 332pp; English.
                                                                                                                                                                                                                                                                                                                                                  19 KAREEEITPUVSIAYKVLEVFPK---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                187 SQTSDWFWCQAANNAN----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR94893 standard; peptide;
                                                                                                                                                                                                                                   Query Match
Best Local Similarity 21.2*
Matches 60; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                 738 AA;
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                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR94893;
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AAR94893
ID AAR9
XX
AC AAR9
XX
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Screening of inhibitors of CD31 is achieved by incubating labelled CD31 component with potential inhibitor, adding this mixture to CD31 component immobilised on a support, washing and detecting label. Alternatively, potential inhibitor and labelled CD31 component immobilised on a support and labelled CD31 component can be added followed by washing and detecting label. Failure to detect label suggests that the compound being screened is not an inhibitor of CD31. The method is used to identify antibodies that can be used in the treatment of carcinomas and inflammation.
                                  domain; antibody; detection; carcinoma; inflammation;
                                                                                                                                                                                                                                                                                  Methods of screening for inhibitors of CD31 interactions
                                                                                                                                                                                                                                                                                             mapping their sites of reaction with the CD31 protein
                                                                                                                                                                                                                                                                                                                    Disclosure, Figure 16, 31pp, English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABG10463 standard; Protein; 506' AA.
                                                                                                                                                                                           (IMCR ) IMPERIAL CANCER RES FUND
          fragment (domains D1-D5)
                                                                                                                                                94GB-0021118
                                                                                                                                                                        94GB-0021118
                                                                                                                                                                                                           (YAMA~) YAMANOUCHI RES INST
                                 CD31; domain; antibodinhibition; treatment
                                                                                                                                                                                                                                  Buckley C,
                                                                                                                                                                                                                                                         WPI; 1996-202498/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      474 AA;
                                                                                                                                              19-OCT-1994;
                                                                        Homo sapiens
                                                                                                                                                                       19-0CT-1994;
                                                                                             GB2294321-A.
                                                                                                                     24-APR-1996.
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          CD31
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diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess New isolated polynucleotide and encoded polypeptides, useful in

Tang YT;

Liu C,

RI,

Drmanac

WPI; 2001-639362/73

N-PSDB; AAS74650

Spragg

Ď,

Simmons

Fawcett J,

30-MAR-2001; 2001WO-US08631.

WO200175067-A2 Homo sapiens.

11-OCT-2001

2000US-0540217.

31-MAR-2000;

23-AUG-2000; 2000US-0649167.

(HYSE-) HYSEQ INC

Claim 20; SEQ ID No 40822; 103pp; English.

biodiversity

9 KVESSRISKVSSIVVNITELFSKPELESSFTHLDQGERINLSCSIPGAPP-----A 325 67 NIKVAKKVVKTHEPASFNLNVTLKSSPDLLTYFCRASSTSGAHVDSARLOMHWELWSKPV 126 SELRANFILQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANFSFLP 186 438 -GRWVLITCCAPQPPPPITYSLCGTK 383 ISYDAQFEVI---KGQTIEVRČESISGTLPISYQLL-KTSKVLENSTKNSNDPAVFKDNP 29; Length 474; 93; Indels 6.9%; Score 97.5; DB 17; 22.8%; Pred. No. 0.94; 27; Mismatches 19 KAREEEITPVVSIAYKVLEVFPK---22.8%; :: : | | | 439 TEDVE-YQCVADN 450 187 SQTSDWFWCQAAN 199 Local Similarity 22.8% nes 44; Conservative

The inventor of the inventor o SELRANFTLODR 138 Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.Wipo.int/pub/published_pct_sequences. 49 VEVPDGITAIWYYDYSGQRRVFMGNPEHRVCNLLLKDLQPEDSGSYNFRFEISEVNRWSD -----KGSCLLIPCIFSFPAD 1 MG-LPGLFCLAVL---AASSFSKAREEEITPVVSIAYKVLEVFPKGRWVLITCCAPQP---GPRVEMICQASSGSPPITNSLIGKDGQVHLQQRP--CHRQPANFS 169 GVKILLSPSGRNILPGELVTLTCQVNSSYPAVSSIKWLKDG-VRLQTKTGVLHLPQAAWS ---GSLVSPPISLHIFMAEVQVS 184 FLPSQTSDWFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLYRSTRRLS Gaps 55 ---PPPIT----YSLCGTKNI-----KVAKKVVKTHEP---ASFNLNVTLKS----97; 6.9%; Score 97.5; DB 22; Length 506; 41; Mismatches 100; Indels The invention relates to isolated polynucleotide (I) 92 -SPDLLTYFCRASSTSGAHVDSARLQMHWELWSKPV-1 MGFLPKLLLLASFFPAGQASWGVSSPQDVQGV-Pred. No. ----DAGVYTCOAENGV---20.7%; 62; Conservative Local Similarity 506 AA; Sequence Query Match à 셤 à d ò 셤 ð d à g Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder.

54 48 243

Novel human diagnostic protein #10454.

(first entry)

13-FEB-2002

Gallatin WM, Vazeux R;

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autoimmunity; inflammation; arthritis; glomerulonephritis;
                                                                                                                                                                                                                                                                                                   label≈ Potential N-glycosylation site
                                                                                                                                                                                                                                                                                                                      'label= Potential N-glycosylation site.
                                                                                                                                                                                                                                                                                                                                           label= Potential N-glycosylation site.
                                                                                                                                                                                                                                                                                                                                                                                  abel= Potential N-glycosylation site.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= Putative transmembrane region.
                                                                                                                                                                                                                                                             /label= Immunoglobulin like domain.
                                                                                                                                                                                                   Immunoglobulin-like domain.
                                                                                                                                                                                                                       'label= Immunoglobulin like domain.
                                                                                                                                                                                                                                          Immunoglobulin like domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                       abel= Potential N-glycosylation
                                                                                                                                                                             /label= putative leader peptide.
43..100
/label= Immunoglobulin-like doma
                                                                                                                                                                                                                                                                               /label= Immunoglobulin like
                                                                                      ICAM-R (Intercellular adhesion molecule R).
                  AAR39741 standard; Protein; 547 AA.
                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-0827689.
92US-0889724.
92US-0894061.
93US-0009266.
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                                                          (updated)
(first entry)
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/label= It
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                                                                                                                      transplant rejection
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                                                                                                                                         Homo sapiens
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26-MAY-1992;
                                                         25-MAR-2003
25-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-JAN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-1992
                                      AAR39741;
                                                                                                          ICAM-R;
                                                                                                                                                            Key
Region
                                                                                                                                                                                         Domain
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RESULT 42
          AAR3974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69 LSKELVASGMGWAAFNLSNVTGNSRILCSVYCNGSQITGSSNITVYGLPERVELAPLPPW 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         123 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    --KNIKVAKKVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9 LWPRACWILLVCCLLTPGVQGQEFILRVEPQNPVLSAGGSLFVNCSTDCPSSEKIALETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLOMHWEL----W
                                                                                                                                                                                    ICAM-R polypeptides can be used in the modulation of immune cell activation/proliferation as competitive inhibitors or stimulatory agents of inter- and intracellular ligand/receptor binding reactions involving ICAM-R. ICAM-R and related products can be used for the treatment of conditions resulting from a response of the non-specific immune response in a mammal. e.g. adult repiratory distress syndrome, acute glomerulonephritis, reactive arthritis, stroke etc, and conditions resulting from a response of the specific immune system in a mammal e.g. psoriasis, or organ/transplant rejection and autoimmune diseases. The ICAM-R products can also be used for monitoring and treating asthma, tumour growth and/or metastasis and viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                55; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6.9%; Score 97.5; DB 14; Length 547; 21.0%; Pred. No. 1.1; ive 41; Mismatches 92; Indels 55.
                                                                                  DNA encoding new human inter-cellular adhesion molecule polypeptide (ICAM-R) - useful for treating immune and inflammatory diseases, tumours and viral infection e.g. HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ---APQPPPPI----TYSLCGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1..29
/label= signal
30..547
/label= ICAM-R
/note= "intercellular adhesion molecule"
                                                                                                                                                      Claim 12; Figure 1(A-G); 126pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human ICAM-R protein.
                                    WPI; 1993-258372/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                          547 AA;
                                                 N-PSDB; AAQ46991
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Best Local S:
Matches 50
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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This sequence represents a novel human intercellular adhesion molecule, ICAM-R. This sequence is used in a method which investigates modulators of the interaction between ICAM-R and the 14.3.3 family member HS1-beta and tubulin. An anti-ICAM-R antibody optionally coupled to toxin or radionuclide, or an ICAM-R aptide, can block; inhibit or stimulate ligand/receptor interactions involving ICAM-R, particularly its effector functions involved in (non)specific immune responses. ICAM-R related agents may be used to treat or monitor inflammation, disorders involving real activation or macrophages, e.g. adult respiratory distress syndrome, stroke, Crohn's disease, multiple sclerosis, rheumatoid arthritis, asthma, tumour growth, human immune deficiency virus infection, diabetes, graft vs. host disease and many others. Antibodies may also be used for passive immunisation, for purifying icam-received to the deficiency of the passive immunisation, for purifying or quantifying ICAM-R expressing cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ---APQPPPPI-----TYSLCGT---KNIKVAKKVV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LWPRACWTLLVCCLLTPGVQGQEFLLRVEPQNPVLSAGGSLFVNCSTDCPSSEKIALETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             55; Gaps
                                                                                                                                                                                                                                                                                                                                                                     Identifying compounds that modulate interaction of intercellular adhesion molecule R - with ligands HSI-beta and tubulin using two-hybrid assay, useful for treating inflammation, T cell activation etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 547;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 1A-G; 108pp; English.
                                                                                                         94US-0286754.
92US-0827689.
92US-0889724.
93US-0809266.
93WG-0102066.
                                                                              95US-0482882
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                                                                                                                                                                                                                                                                                        Vazeux
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                                                                                                                                                                                                                                                          CORP.
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                                                                                                                                                                                                                                                                                                                                       N-PSDB; AAV56348
                                                                                                                                                                                                                                                          SOCI (-SOCI)
                                                                                                                                                                                                                                                                                      Gallatin WM,
                                                                              1995; 1995;
               US5773218-A
                                                                                                                                                         05-JUN-1992
22-JAN-1993
                                             30-JUN-1998
                                                                                                                             27-JAN-1992
                                                                                                                                            26-MAY-1992
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AAW71252 standard; Protein; 547 AA

RESULT 44 AAW71252 AAW71252

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Human; ICAM-R; intercellular adhesion molecule; adhesion; treatment;
inflammatory condition; asthma; tumour growth; metastasis;
viral infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "potential N-linked glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "potential N-linked glycosylation site"
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                                                                                                                                                                                                                                                                                                                                                                                                "putative immunoglobulin-like domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           315..416 // Andrew newnoglobulin-like domain" // Andres "putative immunoglobulin-like domain" // A85 // A85 // A95 // A95
                                                                                                                                                                                                                                                                                                                                                                                                                                           'note= "putative immunoglobulin-like domain"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           486..510*/note= "hydrophobic "transmembrane"
                                                                                  Human intercellular adhesion molecule-R (ICAM-R)
                                                                                                                                                                                                                                                                                                           /note= "signal peptide"
                                                                                                                                                                                                                                                                                                                                                      "mature protein"
                                                                                                                                                                                                                                                                  Location/Qualifiers
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92WS-0827689.
92WS-0889724.
92WS-0894061.
93WS-0102865.
                    (updated)
(first entry)
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N-PSDB; AAV54826.
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05-JUN-1992;
22-JAN-1993;
                    25-MAR-2003
18-NOV-1998
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27-JAN-1992;
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                                                                                                                                                                                                                                                                                    Peptide
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N-PSDB; AAV11657
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26-MAY-1992;
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                                                                                                                                                                    Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents human ICAM-R (intercellular adhesion molecule-R). ICAMs are polypeptides that are expressed on blood vessel endothelial cell surfaces and are involved in the adhesion events in various conditions. ICAM-R variants (see AMW71264-69) can be used to treat or monitor inflammatory conditions involving specific or nonspecific immune responses, asthma, tumour growth and/or metastasis and viral infections. The ICAM variants are produced recombinantly, from expression libraries of mutated sequences, and the ones that are claimed are the ones that have been found to be especially involved in adhesion events. They can also be used to raise antibodies, also for use as
                                                                                                                                                                                                                                                                                                                               123 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPIINSLIGKDGQVHLQQRPCHRQPANF 182
                                                                                                                                                                                                                                                                                                                76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLQMHWEL----W 122
                                                                                                                                                                                                                                                                                                                                                                        183 SFLPSQTSD----WFWCQAANNANVQHSALTVVPPGGDQXMEDWQGPLESPILALPLY 236
                                                                                                                                                                                                                                                                                                                                                                                                                    ICAM-4; ICAM-R; intercellular adhesion molecule; rat; neuron-specific;
promoter; hippocampus; antibody; cell-cell interaction; ss.
                                                                                                                                                                                                                                                                          -----KNIKVAKKVV
                                                                                                                                                                                                                                                                                    DNA encoding mutant ICAM-R poly:peptide(s) - useful for diagnosis and treatment of cell adhesion based disease conditions e.g. inflammation or asthma
                                                                                                                                                                                                                                                      55;
                                                                                                                                                                                                                              / Match 6.9%; Score 97.5; DB 19; Length 547; Local Similarity 21.0%; Pred. No. 1.1; DB 55; Conservative 41; Mismatches 92; Indels 55;
                                                                                                                                                                                                                                                   92; Indels
                                                                                                                                                                        therapeutic or diagnostic agents. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW59005 standard; Protein; 547 AA.
                                         Claim 1; Fig 1A-G; 111pp; English.
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94US-0245295.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                        38 VFPKGRWVLITCC--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ICAM-R protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-311408/27
                                                                                                                                                                                                        547 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SODI (-SODI)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-AUG-1993;
18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5753502-A.
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                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW59005
                                                                                                                                                                                                                                                 Matches
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 D<sub>D</sub>
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11;
                                                                                                                       This sequence represents a human neuron-specific intercellular adhesion molecule, ICAM-R, which is used in a method to isolate a human ICAM-4 gene promoter. This promoter specifically promotes gene transcription in neuronal cells especially hippocampal cells. Recombinant proteins can also be used to raise antibodies against ICAM-4. The ICAM-4 DNA sequences and its recombinant production are new tools in the elucidation of cell-cell interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 123 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rat; intracellular adhesion molecule; ICAM; probe; hybridisation; human; reverse transcription; RT-PCR; RACE; rapid amplification of cDNA ends;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  183 SFLPSQTSD----WFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLY 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ---APQPPPPI----TYSLCGT---KNIKVAKKVV
ICAM-4 gene promoter - for directing gene expression in neuronal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLQMHWEL
                                                                                                                                                                                                                                                                                                                                                                                      Length 547;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                            6.9%; Score 97.5; DB 19; larity 21.0%; Pred. No. 1.1; Conservative 41; Mismatches 92;
                                                                           Disclosure; Col 41-46; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ā
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 38 VFPKGRWVLITCC------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-0245295.
92US-0827689.
92US-0889724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW44838 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-0245295.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human ICAM-4 protein.
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 50; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunogen; antibody
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                                                                                                                                                                                                                                                                                                                                547 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORP.
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us-09-990-726-223.rag

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76 KTHEP-----ASFNINVTLKSSPDILITYFCRASSTSG-AHVDSARLOMHWEL----W 122
                                                                                                                                                                                                                                                                                                                                                                                                                           ICAM; immunoglobulin-like loop; intercellular adhesion molecule receptor; alpha d/CD18; antibody; immunisation; inflammatory response; asthma; tumour growth; viral infection; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         molecule 4 (ICAM-4) protein. The gene sequence was initially isolated from a human P1 library using primers (AAV19345-46) based on the sequence encoding the domain 3 of the human ICAM-3 protein. The screens resulted in the isolation of a 7.0 kb BamHI fragment that hybridised under high stringency conditions with the rat ICAM-4 (AAV1938) sequence. Cells expressing ICAM-4 can be used as immunogens for antibody production and to produce recombinant ICAM-4, which can also be used for antibody production. The DNA can also be used in hybridisation assays to detect expression of ICAM-4 in cells.
                                                                                                                                                                                                                                                                                                                                                                          LWPRACWTLINGCLLTPGVQGQEFLLRVEPONPVLSAGGSLFVNCSTDCPSSEKIALETS 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SFLPSQTSD----WFWCQAANNANVQHSALTVVPPGGDQKWEDWQGPLESPILALPLY 236
                                                                                                                                                                                                                                                                                                                                                 ----APQPPPPI-----TYSLCGT---KNIKVAKKVV
                                                                                                                                                                                                                                                                                                                      55; Gaps
                                                                                   This is the amino acid sequence of the human intracellular adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           174 TATVLASRDDHGAPFSCRIELDMQPQGLGL-FVNTSAPRQLRTFVLPVTPPRLVAPRF
              - useful for transforming cells and producing
                                                                                                                                                                                                                                                                                   6.9%; Score 97.5; DB 19; Length 21.0%; Pred. No. 1.1; Live 41; Mismatches 92; Indels
                                                        Example 9; Column 31-36; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY00779 standard; Protein; 547 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ICAM-R protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-0286754.
92US-0827689.
92US-0889724.
92US-0894061.
93US-0009266.
93WO-US00787.
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95US-0483932.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                               38 VFPKGRWVLITCC--
            DNA encoding ICAM-4
                                                                                                                                                                                                                                                         547 AA;
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                             recombinant
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                                                                                                                                                                                                                                                         Sequence
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Matches
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The invention relates to antibodies (Ab) which bind specifically to the invention relates to antibodies (Ab) which bind specifically to the interaction between ICAM-R and alpha d/CDIB. Abs with specific ICAM-R binding are useful in compositions for immunisation, and for purifying ICAM-R polypeptides and identifying cells expressing ICAM-R on their cell surface, modulating light and biological activities involving ICAM-R, especially inflammatory responses of the specific immune system, the non-specific immune system, monitoring and treating aethma, tumour growth, and/or metassessis, and viral infection (e.g. HIV infection). In particular diseases involving an essential T cell activation (e.g. asthma, psoriasis, diabetes, graft vs. host disease, tissue transplant rejection, and multiple sclerosis) may be treated with anti-ICAM-R antibodies. The Abs specifically bind to and identify ICAM-R and disrupt ICAM-R to cell adhesion molecule, especially alpha d/CDIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLQMHWEL----W 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69 LSKELVASGMGWAAFNLSNVTGNSRILCSVYCNGSQITGSSNITVYGLPERVELAPLPPW 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             123 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ----APOPPPPI----TYSLCGT----KNIKVAKKVV 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Intercellular adhesion molecule polypeptide; ICAM-R; humanised; ICR 1.1; ICR 8.1; monoclonal antibody; therapeutic; inflammatory; asthma; tumour; graft-versus-host disease; viral infection; toxin; radionuclide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    183 SFLPSQTSD----WFWCQAANNANVQHSALTVVPPGGDQKMEDWQGPLESPILALPLY 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ::::: | | | : | | : | 130 | ::: | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | | : | : | | : | : | | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | 
                                                                                                                                                                            New intercellular adhesion molecule receptor (ICAM-R) specific antibodies - useful for modulating ligand/receptor binding and biological activities involving ICAM-R, especially those of the specific and non-specific immune systems
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6.9%; Score 97.5; DB 20; Length 547; 21.0%; Pred. No. 1.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents the human ICAM-R protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "putative leader sequence"
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                                                                                                                                                                                                                                                                                                                                                            Example 5; Fig 1; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW81440 standard; Protein; 547
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        38 VFPKGRWVLITCC---
Vazeux R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neovascularisation site.
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                                                                          WPI; 1999-204041/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       547 AA;
   Gallatin WM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-FEB-1999
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molecule polypeptide (ICAM-R). The invention relates to humanised ICR 1.1 and ICR8 1. antibodies targeted to the ICAM-R polypeptide. Antibodies sand ICR8 8.1 antibodies targeted to the ICAM-R polypeptide. Antibodies specific for ICAM's are potentially useful as therapeutic compounds for treating e.g. immune-mediated inflammatory conditions (e.g. graft-versus-host disease), asthma, tumours or viral infections. Monoclonal antibodies specific for ICAM-R, or their conjugates formed with e.g. toxins or radionuclides are useful for therapeutically targeting or detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-human immunodeficiency virus; HIV; cytostatic; ICAM-R; ARDS; stroke; intercellular adhesion molecule; immunoglobulin heavy chain; septicaemia; inflammatory conditions; glomerulonephritis; arthritis; dermatosis; haemodialysis; leukapheresis, ulcerative colitis; Crohn's disease; necrotising enterocolitis; atherosclerosis; psoriasis, asthma; transplant rejection; diabetes; tumour.
                                                                                                                                                                                                                                                                                                                             SKPVSELRANFTLODRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
                                                                                                                                                                                                                                            ----TYSLCGT---KNIKVAKKVV 75
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                                                                                                                                                                                                                                                                                                          76 KTHEP-----ASFNINVTLKSSPDLLTYFCRASSTSG-AHVDSARLQMHWEL----W
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                                                                                                                                                                           6.9%; Score 97.5; DB 20; 21.0%; Pred. No. 1.1;
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92US-0827689.
92US-0889724.
92US-0894061.
93US-0009266.
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                                                                                                              neovascularisation sites.
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N-PSDB; AAA97090.
                                                                                                                                                                                            Similarity
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22-JAN-1993;
05-AUG-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "putative carboxy terminal cytoplasmic region"
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/note= "putative hydrophobic transmembrane region"
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le - useful for therapeutic or diagnostic purposes
                                                              immunoglobulin-like loop
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                               "potential N-glycosylation site"
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 'note= "mature protein"
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                                                               note= "putative
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92US-0827689.
92US-0889724.
92US-0009266.
93WS-0102862.
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N-PSDB; AAV69125.
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its carboxy terminus. ICAM-R polypeptides are useful for treating and monitoring inflammatory conditions such as adult respiratory distress syndrome, multiple organ injury syndrome secondary to septicaenia or arthritis, dermatosis, stroke, thermal injury, haemodialysis, leukapheresis, ulcerative colitis, Crohn's disease, necrotising artheresis, ulcerative colitis, Crohn's disease, necrotising atheresis, ulcerative colitis, Crohn's disease, necrotising atheresis, ulcerative colitis, Crohn's disease, necrotising atheresis, ulcerating conditions resulting from a response of the also useful for treating conditions resulting from a response of the specific immune system in a mammal e.g. psociasis, organ/tissue transplant rejection and autoimmune diseases including Raymaud's syndrome, autoimmune thyroiditis, multiple sclerosis, rheumatoid arthritis, diabetes and lupus erythematosus. ICAM-R products and ICAM-R calad products are also useful in monitoring and treating asthma, tumour growth and/or metastasis, and viral infection (e.g. HIV infection). Sequences AAA97090 and AAB13036 represent the human ICAM-R DNA and protein sequences. Sequences AAA97112 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primers used in the production of humanised anti-ICAM-R antibody ICR-8.1, and fragments of the humanised antibody. Sequences AAA97124-A97128, AAA97132, AAA97144 represent ICR-8.1 sequences. Sequences AAA97153-A97156 represent primers used in the production of humanised anti-ICAM-R antibody ICR-1.1, and fragments of the humanised antibody. Sequences AAA97155-A97156 and AAB13047-B13048 represent murine
                                                                                                                                                                intercellular adhesion molecuie (ICAM-R) amino acid fragment at its amino terminus and a constant domain of an immunoglobulin heavy chain at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ICR-1.1 sequences. DNA and peptide sequences used in the production of the chimeric protein of the invention include AAA97177-A97188 and
inflammatory conditions, Crohn's
                                                                                                                                  This invention relates to a hybrid fusion protein comprising an
its variants useful, for treating infl
disease, atherosclerosis and diabetes
                                                                                Claim 1; Figure 1; 109pp; English
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547 AA; Sequence

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11;
                                                                                                                                                  76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLOMHWEL----W 122
                                                                                                                                                                                                                                                   -QPVGQ---NFTLR------CQVEGGSPRTSLTVVLLRWEEELSRQPAVEEPAEV 173
                                                                                                                                                                                                                           123 SKPVSELRANFTLQDRGAGPRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182
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                                                                         ----APQPPPPI----TYSLCGT---KNIKVAKKVV 75
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LWPRACWTLLVCCLLTPGVQGQEFLLRVEPQNPVLSAGGSLFVNCSTDCPSSEKIALETS
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 6.9%; Score 97.5; DB 21; Length 547;
                                      Indels
                                   92;
                                    41; Mismatches
                     Pred, No.
                   21.0%;
                                                                         38 VFPKGRWVLITCC----
                                      Conservative
                 Best Local Similarity
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Query Match
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AAY82435 standard; Protein; 547 AA. AAY82435; RESULT 50 AAY82435

(first entry) 28-JUN-2000

Human ICAM-R encoding cDNA SEQ ID NO:1.

Human; ICAM-R; chromosome 19; intracellular adhesion molecule receptor; CAM; ICAM-1; ICAM-2; humanised; antibody; mutagenic; chimeric; vulnerary; nephropathic; antiarthritic; cerebroprotective; antiulcer; cytostatic;

antiarteriosclerotic; immunosuppressive; antidiabetic; neuroprotective; antititytoid; dermatological; antiasthmatic; antibitiammatory; anti-HIV; vasotropic; antipocritic; immunomodulator; antirheumatic; cell adhesion mediator; inflammatory condition; immunisation; immune response

Homo sapiens

US6040176-A.

21-MAR-2000

96US-0714017. 12-SEP-1996;

92US-0827689. 92US-0889724. 92US-0894061. 93US-0009266. 05-AUG-1994 27-JAN-1992 26-MAY-1992

05-JUN-1992

93US-0102852 05-AUG-1993; 22-JAN-1993 26-JAN-1993

(ICOS-) ICOS CORP.

Vazeux Ballatin WM,

WPI; 2000-270138/23.

Novel monoclonal antibody directed against ICAM-R proteins useful for treating acute glomerulonephritis, ulcerative colitis, psoriasis, rheumatoid arthritis, diabetes, multiple sclerosis, asthma and viral Infection

Example 4; Fig 1; 117pp; English.

The present invention describes a monoclonal antibody (MAD) (1), produced by the hybridoma cell line BIKZF (ATCC HB 11692). Also described are: (1) a hybridoma cell line BIKZF, and (2) a MAD (11), that competes with (1) for binding to ICAM-R (intracellular adhesion molecule receptor) (III). (II) mimics the activity of natural binding proteins through which intercellular and intracellular activities of (III) are modulated. (II) is also used for modulating the immune responses. (1) is used for immunisation as well as for purifying (III). They are also used for immunisation as well as for purifying (III). They are also used for immune system in modulating the ligand/receptor binding biological activity involving (III) especially those effector functions of (III) involved in specific and non-specific immune system responses. Inflammatory conditions which may be treated or monitored with related products of (III) include conditions resulting from a response of the non-specific immune system in amammal e.g. adult respiratory distress syndrome, multiple organ injury syndrome secondary to septicemia or trauma, reperfusion injury of tissue, acute glomerulonephritis, reactive arthritis, stroke, ulcerative colitis and atherosclerosis, and conditions the product in mammal, e.g. periatis, organiting from a response of the specific immune system in a mammal, e.g. periatis, organiting from a response of the specific immune system in a mammal, e.g. periatis, organitism transplantation rejection, autoimmune diseases and periatis. such as autoimmune thyroiditis, multiple solerosis, rheumatoid arthritis, diabetes and lupus erythematosus. AAA08236 to AAA08334, and AAY82435 to AAY82451 represent sequences used in the exemplification of the present

Sequence

----APQPPPPI----TYSLCGT---KNIKVAKKVV 75 9 LWPRACWTLLVCCLLTPGVQGQEFLLRVEPQNPVLSAGGSLFVNCSTDCPSSEKIALETS 68 55; Length 547; Indels DB 21; 6.9%; Score 97.5; D 11.0%; Pred. No. 1.1; .ve 41; Mismatches 21.0%; 38 VFPKGRWVLITCC---Conservative Similarity 50; Query Match Local Best Loc Matches

d à

76 KTHEP-----ASFNLNVTLKSSPDLLTYFCRASSTSG-AHVDSARLOMHWEL----W 122

PRVEMICQASSGSPPITNSLIGKDGQVHLQQRPCHRQPANF 182	 CQVEGGSPRTSLTVVLLRWEEELSRQPAVEEPAEV 173
KPVSELRANFTLQDRGA	 OPVGQNFTLR
123	129
ò	qq

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Search completed: February 5, 2004, 16:27:54 Job time: 45 secs